# **Human Secreted Proteins**

This application is a continuation-in-part of, and claims benefit under 35 U.S.C. [1] § 119(e) based on copending U.S. Provisional Application No. 60/278,650 filed on March 27, 2001. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending U.\$. Utility Application No. 09/833,245, filed on April 12, 2001, and PCT International Application Serial No. US01/11988, filed on April 12, 2001. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06043, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/167,061, filed on November 23, 1999, and U.S. Provisional Application No. 60/124,146, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06012, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/166,989, filed on November 23, 1999, and U.S. Provisional Application No. 60/124,093, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06058, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,654, filed on December 3, 1999, and U.S. Provisional Application No. 60/124,145, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06044, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,661, filed on December 3, 1999, and U.S. Provisional Application No. 60/124,099, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT\International Application Serial No. US00/06059, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,622, filed on December 3, 1999, and U.S. Provisional Application No. 60/124,096, filed on March 12, 1999. This application

is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06042, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,663, filed on December 3, 1999, and U.S. Provisional Application No. 60/124,143, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06014, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,665, filed on December 3, 1999, and U.S. Provisional Application No. 60/138,598, filed on June 11, 1999, and U.S. Provisional Application No. 60/124,095, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06013, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 1 19(e) based on U.S. Provisional Application No. 60/168,662, filed on December 3, 1999, and U.S. Provisional Application No. 60/138,626, filed on June 11,1999, and U.S. Provisional Application No. 60/125,360, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06049, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,667, filed on December 3, 1999, and U.S. Provisional Application No. 60/138,574, filed on June 11, 1999, and U.S. Provisional Application No. 60/124,144, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of dopending PCT International Application Serial No. US00/06057, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,666, filed on December 3, 1999, and U.S. Provisional Application No. 60/138,597, filed on June 11, 1999, and U.S. Provisional Application No. 60/124,142, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00 06824, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,664, filed on December 3, 1999, and U.S. Provisional Application No. 60/125,359, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06765, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,906, filed on December 10, 1999, and

U.S. Provisional Application No. 60/126,051, filed on March 23, 1999. This application is also a continuation-in part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06792, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,980, filed on December 10, 1999, and U.S. Provisional Application No. 60/125,362, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06830, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provistonal Application No. 60/169,910, filed on December 10, 1999, and U.S. Provisional Application No. 60/125,361, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06782, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,936, filed on December 10, 1999, and U.S. Provisional Application No. 60/125,812, filed on March 23, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06822, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,916, filed on December 10, 1999, and US. Provisional Application No. 60/126,054, filed on March 23, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06791, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,946, filed on December 10, 1999, and U.S. Provisional Application No. 60/123,815, filed on March 23, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No.\US00/06828, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,616, filed on December 8, 1999, and U.S. Provisional Application No. 60/125,358, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06823, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,623, filed on December 8, 1999, and U.S. Provisional Application No. 60/125,364, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06781, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,617, filed on December 8, 1999, and U.S. Provisional Application No. 60/125,363, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07505, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/172,410, filed on December 17, 1999, and U.S. Provisional Application No. 60/126,502, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07440, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/172,409, filed on December 17, 1999, and U.S. Provisional Application No. 60/126,503, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under §5 U.S.C. § 120 of copending PCT International Application Serial No. US00/07506, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/172,412, filed on December 17, 1999, and U.S. Provisional Application No. 60/126,505, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07507, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/172,408, filed on December 17, 1999, and U.S. Provisional Application No. 60/126,594, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. \$\square\$\text{S00/07535}\$, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/172,413, filed on December 17, 1999, and U.S. Provisional Application No. 60/126,511, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07525, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/171,549, filed on December 22, 1999, and U.S. Provisional Application No. 60/126,595, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07534, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No.

60/171,504, filed on December 22, 1999, and U.S. Provisional Application No. 60/126,598, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07483, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/171,552, filed on December 22, 1999, and U.\$\text{\text{Provisional Application No. 60/126,596, filed on March 26, 1999.} This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07526, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60(171,550, filed on December 22, 1999, and U.S. Provisional Application No. 60/126,600, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07527, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 19(e) based on U.S. Provisional Application No. 60/171,551, filed on December 22, 1999, and U.S. Provisional Application No. 60/126,501, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07661, filed on March 23\, 2000, which claims benefit under 35 U.S.C. \ 119(e) based on U.S. Provisional Application No. 60/174,847, filed on January 7, 2000, and U.S. Provisional Application No. 60/126\504, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. U\$00/07579, filed on March 23, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/174,853, filed on January 7, 2000, and U.S. Provisional Application No. 60/126,509, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07723, filed on March 23, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/242,710, filed on October 25, 2000, and U.S. Provisional Application No. 60/174,850, filed on January 7, 2000, and U.S. Provisional Application No. 60/126,506, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07724, filed on March 23, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/174,850, filed on January 7, 2000, and U.S. Provisional Application No. 60/126,510, filed on

March \$6, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14929, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/174,851, filed on January 7, 2000, and U.S. Provisional Application No. 60/138,573, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07722, filed on March 23, 2000, which claims benefit under 35\U.S.C. § 119(e) based on U.S. Provisional Application No. 60/174,871, filed on January 7, 2000, and U.S. Provisional Application No. 60/126,508, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07578, filed on March 23, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/174,872, filed on January 7, 2000, and U.S. Provisional Application No. 60/126,507, filed on March 26, 1999. This application is also a continuation-in-part of,\and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07726, filed on March 23, 2000, which claims benefit under 35 U.S.C. § 1\19(e) based on U.S. Provisional Application No. 60/174,877, filed on January 7, 2000, and U.S. Provisional Application No. 60/126,597, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07677, filed on March 23\, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/176,064, filed on January 14, 2000, and U.S. Provisional Application No. \( \)0/154,373, filed on September 17, 1999, and U.S. Provisional Application No. 60/126,601, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. U\$00/07725, filed on March 23, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/176,063, filed on January 14, 2000, and U.S. Provisional Application No. 60/126,602, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/09070, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/176,052, filed on January 14, 2000, and U.S. Provisional Application No. 60/128,695, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT

International Application Serial No. US00/08982, filed on April 6, 2000, which claims benefit under \35 U.S.C. \§ 119(e) based on U.S. Provisional Application No. 60/176,069, filed on January 14, 2000, and U.S. Provisional Application No. 60/128,696, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/08983, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/176,068, filed on January 14, 2000, and U.S. Provisional Application No. 60/128,703, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/09067, filed on April 6, 2000, which claims benefit under 35 U.S. & 119(e) based on U.S. Provisional Application No. 60/176,929, filed on January 20, 2000, and U.S. Provisional Application No. 60/128,697, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/09066, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/176,926, and U.S. Provisional Application No. 60/128,698, filed on April 9,\1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.Q. § 120 of copending PCT International Application Serial No. US00/09068, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/177,050, filed on January 20, 2000, and U.S. Provisional Application No. 60/128,699, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial Nd US00/08981, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/177,166, filed on January 20, 2000, and U.S. Provisional Application No. 60/128,701, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/08980, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/176,930, filed on January 20, 2000, and U.S. Provisional Application No. 60/128,700, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/09071, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.\$. Provisional Application No. 60/176,931, filed on January 20, 2000, and U.S. Provisional Application No. 60/128,694, filed on

April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/09069, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/177,049, filed on January 20, 2000, and U.S. Provisional Application No. 60/128,702, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/15136, filed on June 1, 2000, which claims benefit under 35 U.S.C\ § 119(e) based on U.S. Provisional Application No. 60/138,629, filed on June 11, 1999.\ This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14926, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/138,628, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14963, filed on June 1, 2000, which claims benefit under 35 U.S\C. § 119(e) based on U.S. Provisional Application No. 60/138,631, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/15135, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/138,632, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14934, filed on June 1, 2000, which claims benefit under 35 U.S.C.\§ 119(e) based on U.S. Provisional Application No. 60/138,599, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 df copending PCT International Application Serial No. US00/14933, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/138,572, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/15137, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/138,625, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14928, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60(138,633, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of

copending PCT International Application Serial No. US00/14973, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/,138,630, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14964, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/138,627, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/26376, filed on September 26, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/155,808, filed on September 27, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/26371, filed on September 26, 2000, which claims benefit under 35 U.S.C.\§ 119(e) based on U.S. Provisional Application No. 60/155,804, filed on September  $2\frac{1}{2}$ , 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/26324, filed on September 26, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/155,807, filed on September 27, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/26323, filed on September 26, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/155,805, filed on September 27, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/26337, filed on September 26, 2000, which claims benefit under 35 U.S.Q § 119(e) based on U.S. Provisional Application No. 60/155,806, filed on September 27, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US01/13318, filed on April 27, 2001, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/212,142, filed on June 16, 2000, and U.S. Provisional Application No. 60/201,194, filed on May 2, 2000. Each of the above referenced PCT applications were published in the English language. Each of the above referenced priority applications are hereby incorporated by reference in their entireties.

### Field of the Invention

The present invention relates to novel proteins. More specifically, isolated nucleic acid molecules are provided encoding novel polypeptides. Novel polypeptides and antibodies that bind to these polypeptides are provided. Also provided are vectors, host cells, and recombinant and synthetic methods for producing human polynucleotides and/or polypeptides, and antibodies. The invention further relates to diagnostic and therapeutic methods useful for diagnosing, treating, preventing and/or prognosing disorders related to these novel polypeptides. The invention further relates to screening methods for identifying agonists and antagonists of polynucleotides and polypeptides of the invention. The present invention further relates to methods and/or compositions for inhibiting or enhancing the production and function of the polypeptides of the present invention.

## Background of the Invention

- [3] Unlike bacterium, which exist as a single compartment surrounded by a membrane, human cells and other eukaryotes are subdivided by membranes into many functionally distinct compartments. Each membrane-bounded compartment, or organelle, contains different proteins essential for the function of the organelle. The cell uses "sorting signals," which are amino acid motifs located within the protein, to target proteins to particular cellular organelles.
- One type of sorting signal, called a signal sequence, a signal peptide, or a leader sequence, directs a class of proteins to an organelle called the endoplasmic reticulum (ER). The ER separates the membrane-bounded proteins from all other types of proteins. Once localized to the ER, both groups of proteins can be further directed to another organelle called the Golgi apparatus. Here, the Golgi distributes the proteins to vesicles, including secretory vesicles, the cell membrane, lysosomes, and the other organelles.
- [5] Proteins targeted to the ER by a signal sequence can be released into the extracellular space as a secreted protein. For example, vesicles containing secreted proteins can fuse with the cell membrane and release their contents into the extracellular space a process called exocytosis. Exocytosis can occur constitutively or after receipt of a triggering signal. In the latter case, the proteins are stored in secretory vesicles (or

secretory granules) until exocytosis is triggered. Similarly, proteins residing on the cell membrane can also be secreted into the extracellular space by proteolytic cleavage of a "linker" holding the protein to the membrane.

Thus there exists a clear need for identifying and using novel secreted polynucleotides and polypeptides. Identification and sequencing of human genes is a major goal of modern scientific research. For example, by identifying genes and determining their sequences, scientists have been able to make large quantities of valuable human "gene products." These include human insulin, interferon, Factor VIII, tumor necrosis factor, human growth hormone, tissue plasminogen activator, and numerous other compounds. Additionally, knowledge of gene sequences can provide the key to treatment or cure of genetic diseases (such as muscular dystrophy and cystic fibrosis).

### Summary of the Invention

The present invention relates to novel secreted proteins. More specifically, isolated nucleic acid molecules are provided encoding novel secreted polypeptides. Novel polypeptides and antibodies that bind to these polypeptides are provided. Also provided are vectors, host cells, and recombinant and synthetic methods for producing human polynucleotides and/or polypeptides, and antibodies. The invention further relates to diagnostic and therapeutic methods useful for diagnosing, treating, preventing and/or prognosing disorders related to these novel polypeptides. The invention further relates to screening methods for identifying agonists and antagonists of polynucleotides and polypeptides of the invention. The present invention further relates to methods and/or compositions for inhibiting or enhancing the production and function of the polypeptides of the present invention.

### **Detailed Description**

### Polynucleotides and Polypeptides

### **Description of Table 1A**

[8] Table 1A summarizes information concerning certain polypnucleotides and polypeptides of the invention. The first column provides the gene number in the

application for each clone identifier. The second column provides a unique clone identifier, "Clone ID NO:Z", for a cDNA clone related to each contig sequence disclosed in Table 1A. Third column, the cDNA Clones identified in the second column were deposited as indicated in the third column (i.e. by ATCC Deposit Number and deposit date). Some of the deposits contain multiple different clones corresponding to the same gene. In the fourth column, "Vector" refers to the type of vector contained in the corresponding cDNA Clone identified in the second column. In the fifth column, the nucleotide sequence identified as "NT SEQ ID NO:X" was assembled from partially homologous ("overlapping") sequences obtained from the corresponding cDNA clone identified in the second column and, in some cases, from additional related cDNA clones. The overlapping sequences were assembled into a single contiguous sequence of high redundancy (usually three to five overlapping sequences at each nucleotide position), resulting in a final sequence identified as SEQ ID NO:X. In the sixth column, "Total NT Seq." refers to the total number of nucleotides in the contig sequence identified as SEQ ID NO:X." The deposited clone may contain all or most of these sequences, reflected by the nucleotide position indicated as "5' NT of Clone Seq." (seventh column) and the "3' NT of Clone Seq." (eighth column) of SEQ ID NO:X. In the ninth column, the nucleotide position of SEO ID NO:X of the putative start codon (methionine) is identified as "5' NT of Start Codon." Similarly, in column ten, the nucleotide position of SEQ ID NO:X of the predicted signal sequence is identified as "5' NT of First AA of Signal Pep." In the eleventh column, the translated amino acid sequence, beginning with the methionine, is identified as "AA SEQ ID NO:Y," although other reading frames can also be routinely translated using known molecular biology techniques. The polypeptides produced by these alternative open reading frames are specifically contemplated by the present invention.

[9] In the twelfth and thirteenth columns of Table 1A, the first and last amino acid position of SEQ ID NO:Y of the predicted signal peptide is identified as "First AA of Sig Pep" and "Last AA of Sig Pep." In the fourteenth column, the predicted first amino acid position of SEQ ID NO:Y of the secreted portion is identified as "Predicted First AA of Secreted Portion". The amino acid position of SEQ ID NO:Y of the last amino acid encoded by the open reading frame is identified in the fifteenth column as "Last AA of ORF".

- SEQ ID NO:X (where X may be any of the polynucleotide sequences disclosed in the sequence listing) and the translated SEQ ID NO:Y (where Y may be any of the polypeptide sequences disclosed in the sequence listing) are sufficiently accurate and otherwise suitable for a variety of uses well known in the art and described further below. For instance, SEQ ID NO:X is useful for designing nucleic acid hybridization probes that will detect nucleic acid sequences contained in SEQ ID NO:X or the cDNA contained in the deposited clone. These probes will also hybridize to nucleic acid molecules in biological samples, thereby enabling a variety of forensic and diagnostic methods of the invention. Similarly, polypeptides identified from SEQ ID NO:Y may be used, for example, to generate antibodies which bind specifically to proteins containing the polypeptides and the secreted proteins encoded by the cDNA clones identified in Table 1A and/or elsewhere herein
- Nevertheless, DNA sequences generated by sequencing reactions can contain sequencing errors. The errors exist as misidentified nucleotides, or as insertions or deletions of nucleotides in the generated DNA sequence. The erroneously inserted or deleted nucleotides cause frame shifts in the reading frames of the predicted amino acid sequence. In these cases, the predicted amino acid sequence diverges from the actual amino acid sequence, even though the generated DNA sequence may be greater than 99.9% identical to the actual DNA sequence (for example, one base insertion or deletion in an open reading frame of over 1000 bases).
- [12] Accordingly, for those applications requiring precision in the nucleotide sequence or the amino acid sequence, the present invention provides not only the generated nucleotide sequence identified as SEQ ID NO:X, and the predicted translated amino acid sequence identified as SEQ ID NO:Y, but also a sample of plasmid DNA containing a human cDNA of the invention deposited with the ATCC, as set forth in Table 1A. The nucleotide sequence of each deposited plasmid can readily be determined by sequencing the deposited plasmid in accordance with known methods
- [13] The predicted amino acid sequence can then be verified from such deposits. Moreover, the amino acid sequence of the protein encoded by a particular plasmid can also be directly determined by peptide sequencing or by expressing the protein in a suitable host cell containing the deposited human cDNA, collecting the protein, and determining its sequence.

- [14] Also provided in Table 1A is the name of the vector which contains the cDNA plasmid. Each vector is routinely used in the art. The following additional information is provided for convenience.
- [15] Vectors Lambda Zap (U.S. Patent Nos. 5,128,256 and 5,286,636), Uni-Zap XR (U.S. Patent Nos. 5,128, 256 and 5,286,636), Zap Express (U.S. Patent Nos. 5,128,256 and 5,286,636), pBluescript (pBS) (Short, J. M. et al., *Nucleic Acids Res. 16:*7583-7600 (1988); Alting-Mees, M. A. and Short, J. M., *Nucleic Acids Res. 17:*9494 (1989)) and pBK (Alting-Mees, M. A. et al., *Strategies 5:*58-61 (1992)) are commercially available from Stratagene Cloning Systems, Inc., 11011 N. Torrey Pines Road, La Jolla, CA, 92037. pBS contains an ampicillin resistance gene and pBK contains a neomycin resistance gene. Phagemid pBS may be excised from the Lambda Zap and Uni-Zap XR vectors, and phagemid pBK may be excised from the Zap Express vector. Both phagemids may be transformed into *E. coli* strain XL-1 Blue, also available from Stratagene
- Vectors pSport1, pCMVSport 1.0, pCMVSport 2.0 and pCMVSport 3.0, were obtained from Life Technologies, Inc., P. O. Box 6009, Gaithersburg, MD 20897. All Sport vectors contain an ampicillin resistance gene and may be transformed into *E. coli* strain DH10B, also available from Life Technologies. See, for instance, Gruber, C. E., et al., *Focus* 15:59 (1993). Vector lafmid BA (Bento Soares, Columbia University, New York, NY) contains an ampicillin resistance gene and can be transformed into *E. coli* strain XL-1 Blue. Vector pCR<sup>®</sup>2.1, which is available from Invitrogen, 1600 Faraday Avenue, Carlsbad, CA 92008, contains an ampicillin resistance gene and may be transformed into *E. coli* strain DH10B, available from Life Technologies. See, for instance, Clark, J. M., *Nuc. Acids Res.* 16:9677-9686 (1988) and Mead, D. *et al.*, *Bio/Technology* 9: (1991).
- [17] The present invention also relates to the genes corresponding to SEQ ID NO:X, SEQ ID NO:Y, and/or a deposited cDNA (cDNA Clone ID). The corresponding gene can be isolated in accordance with known methods using the sequence information disclosed herein. Such methods include, but are not limited to, preparing probes or primers from the disclosed sequence and identifying or amplifying the corresponding gene from appropriate sources of genomic material.
- [18] Also provided in the present invention are allelic variants, orthologs, and/or species homologs. Procedures known in the art can be used to obtain full-length genes, allelic variants, splice variants, full-length coding portions, orthologs, and/or species homologs of genes corresponding to SEQ ID NO:X and SEQ ID NO:Y using information

from the sequences disclosed herein or the clones deposited with the ATCC. For example, allelic variants and/or species homologs may be isolated and identified by making suitable probes or primers from the sequences provided herein and screening a suitable nucleic acid source for allelic variants and/or the desired homologue.

The present invention provides a polynucleotide comprising, or alternatively consisting of, the nucleic acid sequence of SEQ ID NO:X and/or a cDNA contained in ATCC Deposit No.Z. The present invention also provides a polypeptide comprising, or alternatively, consisting of, the polypeptide sequence of SEQ ID NO:Y, a polypeptide encoded by SEQ ID NO:X, and/or a polypeptide encoded by a cDNA contained in ATCC deposit No.Z. Polynucleotides encoding a polypeptide comprising, or alternatively consisting of the polypeptide sequence of SEQ ID NO:Y, a polypeptide encoded by SEQ ID NO:X and/or a polypeptide encoded by the cDNA contained in ATCC Deposit No.Z, are also encompassed by the invention. The present invention further encompasses a polynucleotide comprising, or alternatively consisting of the complement of the nucleic acid sequence of SEQ ID NO:X, and/or the complement of the coding strand of the cDNA contained in ATCC Deposit No.Z.

### **Description of Table 1B**

Table 1B summarizes some of the polynucleotides encompassed by the [20] invention (including cDNA clones related to the sequences (Clone ID NO:Z), contig sequences (contig identifier (Contig ID:) and contig nucleotide sequence identifier (SEQ ID NO:X)) and further summarizes certain characteristics of these polynucleotides and the The first column provides the gene number in the polypeptides encoded thereby. application for each clone identifier. The second column provides a unique clone identifier, "Clone ID NO:Z", for a cDNA clone related to each contig sequence disclosed in Table 1A and/or 1B. The third column provides a unique contig identifier, "Contig ID:" for each of the contig sequences disclosed in Table 1B. The fourth column provides the sequence identifier, "SEQ ID NO:X", for each of the contig sequences disclosed in Table 1A and/or 1B. The fifth column, "ORF (From-To)", provides the location (i.e., nucleotide position numbers) within the polynucleotide sequence of SEQ ID NO:X that delineate the preferred open reading frame (ORF) that encodes the amino acid sequence shown in the sequence listing and referenced in Table 1B as SEQ ID NO:Y (column 6). Column 7 lists residues comprising predicted epitopes contained in the polypeptides encoded by each of the preferred ORFs (SEQ ID NO:Y). Identification of potential immunogenic regions was performed according to the method of Jameson and Wolf (CABIOS, 4; 181-186 (1988)); specifically, the Genetics Computer Group (GCG) implementation of this algorithm, embodied in the program PEPTIDESTRUCTURE (Wisconsin Package v10.0, Genetics Computer Group (GCG), Madison, Wisc.). This method returns a measure of the probability that a given residue is found on the surface of the protein. Regions where the antigenic index score is greater than 0.9 over at least 6 amino acids are indicated in Table 1B as "Predicted Epitopes". In particular embodiments, polypeptides of the invention comprise, or alternatively consist of, one, two, three, four, five or more of the predicted epitopes described in Table 1B. It will be appreciated that depending on the analytical criteria used to predict antigenic determinants, the exact address of the determinant may vary slightly. Column 8, "Tissue Distribution" shows the expression profile of tissue, cells, and/or cell line libraries which express the polynucleotides of the invention. The first number in column 8 (preceding the colon), represents the tissue/cell source identifier code corresponding to the key provided in Table 4. Expression of these polynucleotides was not observed in the other tissues and/or cell libraries tested. For those identifier codes in which the first two letters are not "AR", the second number in column 8 (following the colon), represents the number of times a sequence corresponding to the reference polynucleotide sequence (e.g., SEQ ID NO:X) was identified in the tissue/cell source. Those tissue/cell source identifier codes in which the first two letters are "AR" designate information generated using DNA array Utilizing this technology, cDNAs were amplified by PCR and then technology. transferred, in duplicate, onto the array. Gene expression was assayed through hybridization of first strand cDNA probes to the DNA array. cDNA probes were generated from total RNA extracted from a variety of different tissues and cell lines. Probe synthesis was performed in the presence of <sup>33</sup>P dCTP, using oligo(dT) to prime reverse transcription. After hybridization, high stringency washing conditions were employed to remove non-specific hybrids from the array. The remaining signal, emanating from each gene target, was measured using a Phosphorimager. Gene expression was reported as Phosphor Stimulating Luminescence (PSL) which reflects the level of phosphor signal generated from the probe hybridized to each of the gene targets represented on the array. A local background signal subtraction was performed before the total signal generated from each array was used to normalize gene expression between the different hybridizations. The value presented after "[array code]:" represents the mean of the duplicate values, following background subtraction and probe normalization. One of skill in the art could routinely use this information to identify normal and/or diseased tissue(s) which show a predominant expression pattern of the corresponding polynucleotide of the invention or to identify polynucleotides which show predominant and/or specific tissue and/or cell expression. Column 9 provides the chromosomal location of polynucleotides corresponding to SEQ ID NO:X. Chromosomal location was determined by finding exact matches to EST and cDNA sequences contained in the NCBI (National Center for Biotechnology Information) UniGene database. Given a presumptive chromosomal location, disease locus association was determined by comparison with the Morbid Map, derived from Online Mendelian Inheritance in Man (Online Mendelian Inheritance in Man, OMIM<sup>TM</sup>. McKusick-Nathans Institute for Genetic Medicine, Johns Hopkins University (Baltimore, MD) and National Center for Biotechnology Information, National MD) 2000. World Wide Web URL: (Bethesda, Library Medicine http://www.ncbi.nlm.nih.gov/omim/). If the putative chromosomal location of the Query overlaps with the chromosomal location of a Morbid Map entry, an OMIM identification number is disclosed in column 10 labeled "OMIM Disease Reference(s)". A key to the OMIM reference identification numbers is provided in Table 5.

### **Description of Table 1C**

Table 1C summarizes additional polynucleotides encompassed by the invention [21] (including cDNA clones related to the sequences (Clone ID NO:Z), contig sequences (contig identifier (Contig ID:) contig nucleotide sequence identifiers (SEQ ID NO:X)), and genomic sequences (SEQ ID NO:B). The first column provides a unique clone identifier, "Clone ID NO:Z", for a cDNA clone related to each contig sequence. The second column provides the sequence identifier, "SEQ ID NO:X", for each contig sequence. The third column provides a unique contig identifier, "Contig ID:" for each contig sequence. The fourth column, provides a BAC identifier "BAC ID NO:A" for the BAC clone referenced in the corresponding row of the table. The fifth column provides the nucleotide sequence identifier, "SEQ ID NO:B" for a fragment of the BAC clone identified in column four of the corresponding row of the table. The sixth column, "Exon From-To", provides the location (i.e., nucleotide position numbers) within the polynucleotide sequence of SEQ ID NO:B which delineate certain polynucleotides of the invention that are also exemplary members of polynucleotide sequences that encode polypeptides of the invention (e.g., polypeptides containing amino acid sequences encoded by the polynucleotide sequences delineated in column six, and fragments and variants thereof).

### **Description of Table 1D**

Table 1D: In preferred embodiments, the present invention encompasses a method of treating a disease or disorder listed in the "FEATURES OF PROTEIN" sections (below) and also as listed in the "Preferred Indications" column of Table 1D (below); comprising administering to a patient in which such treatment, prevention, or amelioration is desired a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) represented by Table 1A and Table 1D (in the same row as the disease or disorder to be treated is listed in the "Preferred Indications" column of Table 1D) in an amount effective to treat, prevent, or ameliorate the disease or disorder.

[23] As indicated in Table 1D, the polynucleotides, polypeptides, agonists, or antagonists of the present invention (including antibodies) can be used in assays to test for one or more biological activities. If these polynucleotides and polypeptides do exhibit activity in a particular assay, it is likely that these molecules may be involved in the diseases associated with the biological activity. Thus, the polynucleotides or polypeptides, or agonists or antagonists thereof (including antibodies) could be used to treat the associated disease.

The present invention encompasses methods of preventing, treating, diagnosing, or ameliorating a disease or disorder. In preferred embodiments, the present invention encompasses a method of treating a disease or disorder listed in the "Preferred Indications" column of Table 1D; comprising administering to a patient in which such treatment, prevention, or amelioration is desired a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) in an amount effective to treat, prevent, diagnose, or ameliorate the disease or disorder. The first and seccond columns of Table 1D show the "Gene No." and "cDNA Clone ID No.", respectively, indicating certain nucleic acids and proteins (or antibodies against the same) of the invention (including polynucleotide, polypeptide, and antibody fragments or variants thereof) that may be used in preventing, treating, diagnosing, or ameliorating the disease(s) or disorder(s) indicated in the corresponding row in Column 3 of Table 1D.

[25] In another embodiment, the present invention also encompasses methods of preventing, treating, diagnosing, or ameliorating a disease or disorder listed in the "Preferred Indications" column of Table 1D; comprising administering to a patient

combinations of the proteins, nucleic acids, or antibodies of the invention (or fragments or variants thereof), sharing similar indications as shown in the corresponding rows in Column 3 of Table 1D.

[26] The "Preferred Indication" column describes diseases, disorders, and/or conditions that may be treated, prevented, diagnosed, or ameliorated by a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof).

The recitation of "Cancer" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof) may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., leukemias, cancers, and/or as described below under "Hyperproliferative Disorders").

[28] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Cancer" recitation in the "Preferred Indication" column of Table 1D may be used for example, to diagnose, treat, prevent, and/or ameliorate a neoplasm located in a tissue selected from the group consisting of: colon, abdomen, bone, breast, digestive system, liver, pancreas, prostate, peritoneum, lung, blood (e.g., leukemia), endocrine glands (adrenal, parathyroid, pituitary, testicles, ovary, thymus, thyroid), uterus, eye, head and neck, nervous (central and peripheral), lymphatic system, pelvic, skin, soft tissue, spleen, thoracic, and urogenital.

In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Cancer" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a pre-neoplastic condition, selected from the group consisting of: hyperplasia (e.g., endometrial hyperplasia and/or as described in the section entitled "Hyperproliferative Disorders"), metaplasia (e.g., connective tissue metaplasia, atypical metaplasia, and/or as described in the section entitled "Hyperproliferative Disorders"), and/or dysplasia (e.g., cervical dysplasia, and bronchopulmonary dysplasia).

[30] In another specific embodiment, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Cancer" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a benign dysproliferative disorder selected from the group consisting of: benign tumors, fibrocystic conditions, tissue hypertrophy, and/or as described in the section entitled "Hyperproliferative Disorders".

- [31] The recitation of "Immune/Hematopoietic" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), blood disorders (e.g., as described below under "Immune Activity" "Cardiovascular Disorders" and/or "Blood-Related Disorders"), and infections (e.g., as described below under "Infectious Disease").
- In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having the "Immune/Hematopoietic" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: anemia, pancytopenia, leukopenia, thrombocytopenia, leukemias, Hodgkin's disease, non-Hodgkin's lymphoma, acute lymphocytic anemia (ALL), plasmacytomas, multiple myeloma, Burkitt's lymphoma, arthritis, asthma, AIDS, autoimmune disease, rheumatoid arthritis, granulomatous disease, immune deficiency, inflammatory bowel disease, sepsis, neutropenia, neutrophilia, psoriasis, immune reactions to transplanted organs and tissues, systemic lupus erythematosis, hemophilia, hypercoagulation, diabetes mellitus, endocarditis, meningitis, Lyme Disease, and allergies.
- [33] The recitation of "Reproductive" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), and disorders of the reproductive system (e.g., as described below under "Reproductive System Disorders").
- [34] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Reproductive" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: cryptorchism, prostatitis, inguinal hernia, varicocele, leydig cell tumors, verrucous carcinoma, prostatitis, malacoplakia, Peyronie's disease, penile carcinoma, squamous cell hyperplasia, dysmenorrhea, ovarian adenocarcinoma, Turner's syndrome, mucopurulent cervicitis, Sertoli-leydig tumors, ovarian cancer, uterine cancer, pelvic inflammatory disease, testicular cancer, prostate cancer, Klinefelter's syndrome, Young's syndrome,

premature ejaculation, diabetes mellitus, cystic fibrosis, Kartagener's syndrome, testicular atrophy, testicular feminization, anorchia, ectopic testis, epididymitis, orchitis, gonorrhea, syphilis, testicular torsion, vasitis nodosa, germ cell tumors, stromal tumors, dysmenorrhea, retroverted uterus, endometriosis, fibroids, adenomyosis, anovulatory bleeding, amenorrhea, Cushing's syndrome, hydatidiform moles, Asherman's syndrome, premature menopause, precocious puberty, uterine polyps, dysfunctional uterine bleeding, cervicitis, chronic cervicitis, mucopurulent cervicitis, cervical dysplasia, cervical polyps, Nabothian cysts, cervical erosion, cervical incompetence, cervical neoplasms, pseudohermaphroditism, and premenstrual syndrome.

- [35] The recitation of "Musculoskeletal" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), and disorders of the immune system (e.g., as described below under "Immune Activity").
- [36] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Musculoskeletal" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: bone cancers (e.g., osteochondromas, benign chondromas, chondroblastoma, chondromyxoid fibromas, osteoid osteomas, giant cell tumors, multiple myeloma, osteosarcomas), Paget's Disease, rheumatoid arthritis, systemic lupus erythematosus, osteomyelitis, Lyme Disease, gout, bursitis, tendonitis, osteoporosis, osteoarthritis, muscular dystrophy, mitochondrial myopathy, cachexia, and multiple sclerosis.
- [37] The recitation of "Cardiovascular" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), and disorders of the cardiovascular system (e.g., as described below under "Cardiovascular Disorders").
- [38] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Cardiovascular" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: myxomas,

fibromas, rhabdomyomas, cardiovascular abnormalities (e.g., congenital heart defects, cerebral arteriovenous malformations, septal defects), heart disease (e.g., heart failure, congestive heart disease, arrhythmia, tachycardia, fibrillation, pericardial Disease, endocarditis), cardiac arrest, heart valve disease (e.g., stenosis, regurgitation, prolapse), vascular disease (e.g., hypertension, coronary artery disease, angina, aneurysm, arteriosclerosis, peripheral vascular disease), hyponatremia, hypernatremia, hypokalemia, and hyperkalemia.

- [39] The recitation of "Mixed Fetal" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders").
- In specific embodiments, a protein, nucleic acid, or antibody of the invention [40] (or fragment or variant thereof) having a "Mixed Fetal" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: spina bifida, hydranencephaly, neurofibromatosis, fetal alcohol syndrome, diabetes mellitus, PKU, Down's syndrome, Patau syndrome, Edwards syndrome, Turner syndrome, Apert syndrome, Carpenter syndrome, Conradi syndrome, Crouzon syndrome, cutis laxa, Cornelia de Lange syndrome, Ellis-van Creveld syndrome, Holt-Oram syndrome, Kartagener syndrome, Meckel-Gruber syndrome, Noonan syndrome, Pallister-Hall syndrome, Rubinstein-Taybi syndrome, Scimitar syndrome, Smith-Lemli-Opitz syndrome, thromocytopenia-absent radius (TAR) syndrome, Treacher Collins syndrome, Williams syndrome, Hirschsprung's disease, Meckel's diverticulum, polycystic kidney disease, Turner's syndrome, and gonadal dysgenesis, Klippel-Feil syndrome, Ostogenesis imperfecta, muscular dystrophy, Tay-Sachs disease, Wilm's tumor, neuroblastoma, and retinoblastoma.
- [41] The recitation of "Excretory" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and renal disorders (e.g., as described below under "Renal Disorders").

- [42] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Excretory" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: bladder cancer, prostate cancer, benign prostatic hyperplasia, bladder disorders (e.g., urinary incontinence, urinary retention, urinary obstruction, urinary tract Infections, interstitial cystitis, prostatitis, neurogenic bladder, hematuria), renal disorders (e.g., hydronephrosis, proteinuria, renal failure, pyelonephritis, urolithiasis, reflux nephropathy, and unilateral obstructive uropathy).
- [43] The recitation of "Neural/Sensory" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and diseases or disorders of the nervous system (e.g., as described below under "Neural Activity and Neurological Diseases").
- In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Neural/Sensory" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: brain cancer (e.g., brain stem glioma, brain tumors, central nervous system (Primary) lymphoma, central nervous system lymphoma, cerebellar astrocytoma, and cerebral astrocytoma, neurodegenerative disorders (e.g., Alzheimer's Disease, Creutzfeldt-Jakob Disease, Parkinson's Disease, and Idiopathic Presenile Dementia), encephalomyelitis, cerebral malaria, meningitis, metabolic brain diseases (e.g., phenylketonuria and pyruvate carboxylase deficiency), cerebellar ataxia, ataxia telangiectasia, and AIDS Dementia Complex, schizophrenia, attention deficit disorder, hyperactive attention deficit disorder, autism, and obsessive compulsive disorders.
- [45] The recitation of "Respiratory" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and diseases or disorders of the respiratory system (e.g., as described below under "Respiratory Disorders").

In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Respiratory" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: cancers of the respiratory system such as larynx cancer, pharynx cancer, trachea cancer, epiglottis cancer, lung cancer, squamous cell carcinomas, small cell (oat cell) carcinomas, large cell carcinomas, and adenocarcinomas. Allergic reactions, cystic fibrosis, sarcoidosis, histiocytosis X, infiltrative lung diseases (e.g., pulmonary fibrosis and lymphoid interstitial pneumonia), obstructive airway diseases (e.g., asthma, emphysema, chronic or acute bronchitis), occupational lung diseases (e.g., silicosis and asbestosis), pneumonia, and pleurisy.

The recitation of "Endocrine" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and diseases or disorders of the respiratory system (e.g., as described below under "Respiratory Disorders"), renal disorders (e.g., as described below under "Renal Disorders"), and disorders of the endocrine system (e.g., as described below under "Endocrine Disorders".

[48] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having an "Endocrine" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: cancers of endocrine tissues and organs (e.g., cancers of the hypothalamus, pituitary gland, thyroid gland, parathyroid glands, pancreas, adrenal glands, ovaries, and testes), diabetes (e.g., diabetes insipidus, type I and type II diabetes mellitus), obesity, disorders related to pituitary glands (e.g., hyperpituitarism, hypopituitarism, and pituitary dwarfism), hypothyroidism, hyperthyroidism, goiter, reproductive disorders (e.g. male and female infertility), disorders related to adrenal glands (e.g., Addison's Disease, corticosteroid deficiency, and Cushing's Syndrome), kidney cancer (e.g., hypernephroma, transitional cell cancer, and Wilm's tumor), diabetic nephropathy, interstitial nephritis, polycystic kidney disease, glomerulonephritis (e.g., IgM mesangial proliferative glomerulonephritis and glomerulonephritis caused by autoimmune disorders; such as Goodpasture's syndrome), and nephrocalcinosis.

- [49] The recitation of "Digestive" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and diseases or disorders of the gastrointestinal system (e.g., as described below under "Gastrointestinal Disorders".
- In specific embodiments, a protein, nucleic acid, or antibody of the invention [50] (or fragment or variant thereof) having a "Digestive" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: ulcerative colitis, appendicitis, Crohn's disease, hepatitis, hepatic encephalopathy, portal hypertension, cholelithiasis, cancer of the digestive system (e.g., biliary tract cancer, stomach cancer, colon cancer, gastric cancer, pancreatic cancer, cancer of the bile duct, tumors of the colon (e.g., polyps or cancers), and cirrhosis), pancreatitis, ulcerative disease, pyloric stenosis, gastroenteritis, gastritis, gastric atropy, benign tumors of the duodenum, distension, irritable bowel syndrome, malabsorption, congenital disorders of the small intestine, bacterial and parasitic infection, megacolon, Hirschsprung's disease, aganglionic megacolon, acquired megacolon, colitis, anorectal disorders (e.g., anal fistulas, hemorrhoids), congenital disorders of the liver (e.g., Wilson's disease, hemochromatosis, cystic fibrosis, biliary atresia, and alpha1-antitrypsin deficiency), portal hypertension, cholelithiasis, and jaundice.
- [51] The recitation of "Connective/Epithelial" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), cellular and genetic abnormalities (e.g., as described below under "Diseases at the Cellular Level "), angiogenesis (e.g., as described below under "Anti-Angiogenesis Activity "), and or to promote or inhibit regeneration (e.g., as described below under "Regeneration "), and wound healing (e.g., as described below under "Wound Healing and Epithelial Cell Proliferation").
- [52] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Connective/Epithelial" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of:

metaplasia, mixed connective tissue disease, focal epithelial hyperplasia, epithelial metaplasia, mucoepithelial dysplasia, graft v. host disease, polymyositis, cystic hyperplasia, cerebral dysplasia, tissue hypertrophy, Alzheimer's disease, lymphoproliferative disorder, Waldenstron's macroglobulinemia, Crohn's disease, pernicious anemia, idiopathic Addison's disease, glomerulonephritis, bullous pemphigoid, Sjogren's syndrome, diabetes mellitus, cystic fibrosis, osteoblastoma, osteoclastoma, osteosarcoma, chondrosarcoma, osteoporosis, osteocarthritis, periodontal disease, wound healing, relapsing polychondritis, vasculitis, polyarteritis nodosa, Wegener's granulomatosis, cellulitis, rheumatoid arthritis, psoriatic arthritis, discoid lupus erythematosus, systemic lupus erythematosus, scleroderma, CREST syndrome, Sjogren's syndrome, polymyositis, dermatomyositis, mixed connective tissue disease, relapsing polychondritis, vasculitis, Henoch-Schonlein syndrome, erythema nodosum, polyarteritis nodosa, temporal (giant cell) arteritis, Takayasu's arteritis, Wegener's granulomatosis, Reiter's syndrome, Behcet's syndrome, ankylosing spondylitis, cellulitis, keloids, Ehler Danlos syndrome, Marfan syndrome, pseudoxantoma elasticum, osteogenese imperfecta, chondrodysplasias, epidermolysis bullosa, Alport syndrome, and cutis laxa.

### **Description of Table 1E**

Table 1E provides information related to biological activities and preferred [53] indications for polynucleotides and polypeptides of the invention (including antibodies, agonists, and/or antagonists thereof). Table 1E also provides information related to assays which may be used to test polynucleotides and polypeptides of the invention (including antibodies, agonists, and/or antagonists thereof) for the corresponding biological activities. The first column ("Gene No.") provides the gene number in the application for each clone identifier. The second column ("cDNA Clone ID No:Z") provides the unique clone identifier for each clone as previously described and indicated in Tables 1A, 1B, 1C, and 1D. The third column ("AA SEQ ID NO:Y") indicates the Sequence Listing SEQ ID Number for polypeptide sequences encoded by the corresponding cDNA clones (also as indicated in Tables 1A, 1B, and 2). The fourth column ("Biological Activity") indicates a biological activity corresponding to the indicated polypeptides (or polynucleotides encoding said The fifth column ("Exemplary Activity Assay") further describes the polypeptides). corresponding biological activity and provides information pertaining to the various types of assays which may be performed to test, demonstrate, or quantify the corresponding biological activity. The sixth column ("Preferred Indications") describes particular embodiments of the invention and indications (e.g. pathologies, diseases, disorders, abnormalities, etc.) for which polynucleotides and polypeptides of the invention (including antibodies, agonists, and/or antagonists thereof) may be used in detecting, diagnosing, preventing, and/or treating.

Table 1E describes the use of FMAT technology, inter alia, for testing or [54] demonstrating various biological activities. Fluorometric microvolume assay technology (FMAT) is a fluorescence-based system which provides a means to perform nonradioactive cell- and bead-based assays to detect activation of cell signal transduction pathways. This technology was designed specifically for ligand binding and immunological assays. Using this technology, fluorescent cells or beads at the bottom of the well are detected as localized areas of concentrated fluorescence using a data processing system. Unbound flurophore comprising the background signal is ignored, allowing for a wide variety of homogeneous assays. FMAT technology may be used for peptide ligand binding assays, immunofluorescence, apoptosis, cytotoxicity, and beadbased immunocapture assays. See, Miraglia S et. al., "Homogeneous cell and bead based assays for highthroughput screening using flourometric microvolume assay technology," Journal of Biomolecular Screening; 4:193-204 (1999). In particular, FMAT technology may be used to test, confirm, and/or identify the ability of polypeptides (including polypeptide fragments and variants) to activate signal transduction pathways. example, FMAT technology may be used to test, confirm, and/or identify the ability of polypeptides to upregulate production of immunomodulatory proteins (such as, for example, interleukins, GM-CSF, Rantes, and Tumor Necrosis factors, as well as other cellular regulators (e.g. insulin)).

Table 1E also describes the use of kinase assays for testing, demonstrating, or quantifying biological activity. In this regard, the phosphorylation and dephosphorylation of specific amino acid residues (e.g. Tyrosine, Serine, Threonine) on cell-signal transduction proteins provides a fast, reversible means for activation and deactivation of cellular signal transduction pathways. Moreover, cell signal transduction via phosphorylation/de-phosphorylation is crucial to the regulation of a wide variety of cellular processes (e.g. proliferation, differentiation, migration, apoptosis, etc.). Accordingly, kinase assays provide a powerful tool useful for testing, confirming, and/or identifying polypeptides (including polypeptide fragments and variants) that mediate cell signal transduction events via protein phosphorylation. See e.g., Forrer, P., Tamaskovic R., and Jaussi, R. "Enzyme-Linked Immunosorbent Assay for Measurement of JNK, ERK, and p38 Kinase Activities" Biol. Chem. 379(8-9): 1101-1110 (1998).

### **Description of Table 2**

Table 2 summarizes homology and features of some of the polypeptides of the [56] invention. The first column provides a unique clone identifier, "Clone ID NO:Z", corresponding to a cDNA clone disclosed in Table 1A or 1B. The second column provides the unique contig identifier, "Contig ID:" corresponding to contigs in Table 1B and allowing for correlation with the information in Table 1B. The third column provides the sequence identifier, "SEQ ID NO:X", for the contig polynucleotide sequence. The fourth column provides the analysis method by which the homology/identity disclosed in the Table was determined. Comparisons were made between polypeptides encoded by the polynucleotides of the invention and either a non-redundant protein database (herein referred to as "NR"), or a database of protein families (herein referred to as "PFAM") as further described below. The fifth column provides a description of the PFAM/NR hit having a significant match to a polypeptide of the invention. Column six provides the accession number of the PFAM/NR hit disclosed in the fifth column. Column seven, "Score/Percent Identity", provides a quality score or the percent identity, of the hit disclosed in columns five and six. Columns 8 and 9, "NT From" and "NT To" respectively, delineate the polynucleotides in "SEQ ID NO:X" that encode a polypeptide having a significant match to the PFAM/NR database as disclosed in the fifth and sixth In specific embodiments polypeptides of the invention comprise, or columns. alternatively consist of, an amino acid sequence encoded by a polynucleotide in SEQ ID NO:X as delineated in columns 8 and 9, or fragments or variants thereof.

### **Description of Table 3**

Table 3 provides polynucleotide sequences that may be disclaimed according to certain embodiments of the invention. The first column provides a unique clone identifier, "Clone ID", for a cDNA clone related to contig sequences disclosed in Table 1B. The second column provides the sequence identifier, "SEQ ID NO:X", for contig sequences disclosed in Table 1A and/or 1B. The third column provides the unique contig identifier,

"Contig ID:", for contigs disclosed in Table 1B. The fourth column provides a unique integer 'a' where 'a' is any integer between 1 and the final nucleotide minus 15 of SEQ ID NO:X, and the fifth column provides a unique integer 'b' where 'b' is any integer between 15 and the final nucleotide of SEQ ID NO:X, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:X, and where b is greater than or equal to a + 14. For each of the polynucleotides shown as SEQ ID NO:X, the uniquely defined integers can be substituted into the general formula of a-b, and used to describe polynucleotides which may be preferably excluded from the invention. In certain embodiments, preferably excluded from the invention are at least one, two, three, four, five, ten, or more of the polynucleotide sequence(s) having the accession number(s) disclosed in the sixth column of this Table (including for example, published sequence in connection with a particular BAC clone). In further embodiments, preferably excluded from the invention are the specific polynucleotide sequence(s) contained in the clones corresponding to at least one, two, three, four, five, ten, or more of the available material having the accession numbers identified in the sixth column of this Table (including for example, the actual sequence contained in an identified BAC clone).

### **Description of Table 4**

Table 4 provides a key to the tissue/cell source identifier code disclosed in Table 1B, column 8. Column 1 provides the tissue/cell source identifier code disclosed in Table 1B, Column 8. Columns 2-5 provide a description of the tissue or cell source. Codes corresponding to diseased tissues are indicated in column 6 with the word "disease". The use of the word "disease" in column 6 is non-limiting. The tissue or cell source may be specific (e.g. a neoplasm), or may be disease-associated (e.g., a tissue sample from a normal portion of a diseased organ). Furthermore, tissues and/or cells lacking the "disease" designation may still be derived from sources directly or indirectly involved in a disease state or disorder, and therefore may have a further utility in that disease state or disorder. In numerous cases where the tissue/cell source is a library, column 7 identifies the vector used to generate the library.

### **Description of Table 5**

[59] Table 5 provides a key to the OMIM reference identification numbers disclosed in Table 1B, column 10. OMIM reference identification numbers (Column 1) were derived from Online Mendelian Inheritance in Man (Online Mendelian Inheritance in

Man, OMIM. McKusick-Nathans Institute for Genetic Medicine, Johns Hopkins University (Baltimore, MD) and National Center for Biotechnology Information, National Library of Medicine, (Bethesda, MD) 2000. World Wide Web URL: http://www.ncbi.nlm.nih.gov/omim/). Column 2 provides diseases associated with the cytologic band disclosed in Table 1B, column 9, as determined using the Morbid Map database.

### **Description of Table 6**

[60] Table 6 summarizes some of the ATCC Deposits, Deposit dates, and ATCC designation numbers of deposits made with the ATCC in connection with the present application. These deposits were made in addition to those described in the Table 1A.

### **Description of Table 7**

- [61] Table 7 shows the cDNA libraries sequenced, and ATCC designation numbers and vector information relating to these cDNA libraries.
- [62] The first column shows the first four letters indicating the Library from which each library clone was derived. The second column indicates the catalogued tissue description for the corresponding libraries. The third column indicates the vector containing the corresponding clones. The fourth column shows the ATCC deposit designation for each library clone as indicated by the deposit information in Table 6.

### **Definitions**

- [63] The following definitions are provided to facilitate understanding of certain terms used throughout this specification.
- [64] In the present invention, "isolated" refers to material removed from its original environment (e.g., the natural environment if it is naturally occurring), and thus is altered "by the hand of man" from its natural state. For example, an isolated polynucleotide could be part of a vector or a composition of matter, or could be contained within a cell, and still be "isolated" because that vector, composition of matter, or particular cell is not the original environment of the polynucleotide. The term "isolated" does not refer to genomic or cDNA libraries, whole cell total or mRNA preparations, genomic DNA preparations

(including those separated by electrophoresis and transferred onto blots), sheared whole cell genomic DNA preparations or other compositions where the art demonstrates no distinguishing features of the polynucleotide/sequences of the present invention.

- In the present invention, a "secreted" protein refers to those proteins capable of being directed to the ER, secretory vesicles, or the extracellular space as a result of a signal sequence, as well as those proteins released into the extracellular space without necessarily containing a signal sequence. If the secreted protein is released into the extracellular space, the secreted protein can undergo extracellular processing to produce a "mature" protein. Release into the extracellular space can occur by many mechanisms, including exocytosis and proteolytic cleavage.
- As used herein, a "polynucleotide" refers to a molecule having a nucleic acid [66] sequence encoding SEQ ID NO:Y or a fragment or variant thereof (e.g., the polypeptide delinated in columns fourteen and fifteen of Table 1A); a nucleic acid sequence contained in SEQ ID NO:X (as described in column 5 of Table 1A and/or column 3 of Table 1B) or the complement thereof; a cDNA sequence contained in Clone ID NO:Z (as described in column 2 of Table 1A and/or 1B and contained within a library deposited with the ATCC); a nucleotide sequence encoding the polypeptide encoded by a nucleotide sequence in SEQ ID NO:B as defined in column 6 (EXON From-To) of Table 1C or a fragment or variant thereof; or a nucleotide coding sequence in SEQ ID NO:B as defined in column 6 of Table 1C or the complement thereof. For example, the polynucleotide can contain the nucleotide sequence of the full length cDNA sequence, including the 5' and 3' untranslated sequences, the coding region, as well as fragments, epitopes, domains, and variants of the nucleic acid sequence. Moreover, as used herein, a "polypeptide" refers to a molecule having an amino acid sequence encoded by a polynucleotide of the invention as broadly defined (obviously excluding poly-Phenylalanine or poly-Lysine peptide sequences which result from translation of a polyA tail of a sequence corresponding to a cDNA).
- [67] In the present invention, "SEQ ID NO:X" was often generated by overlapping sequences contained in multiple clones (contig analysis). A representative clone containing all or most of the sequence for SEQ ID NO:X is deposited at Human Genome Sciences, Inc. (HGS) in a catalogued and archived library. As shown, for example, in

column 2 of Table 1B, each clone is identified by a cDNA Clone ID (identifier generally referred to herein as Clone ID NO:Z). Each Clone ID is unique to an individual clone and the Clone ID is all the information needed to retrieve a given clone from the HGS library. Table 7 provides a list of the deposited cDNA libraries. One can use the Clone ID NO:Z to determine the library source by reference to Tables 6 and 7. Table 7 lists the deposited cDNA libraries by name and links each library to an ATCC Deposit. Library names contain four characters, for example, "HTWE." The name of a cDNA clone (Clone ID) isolated from that library begins with the same four characters, for example "HTWEP07". As mentioned below, Table 1A and/or 1B correlates the Clone ID names with SEQ ID NO:X. Thus, starting with an SEQ ID NO:X, one can use Tables 1A, 1B, 6, 7, and 9 to determine the corresponding Clone ID, which library it came from and which ATCC deposit the library is contained in. Furthermore, it is possible to retrieve a given cDNA clone from the source library by techniques known in the art and described elsewhere herein. The ATCC is located at 10801 University Boulevard, Manassas, Virginia 20110-2209, USA. The ATCC deposits were made pursuant to the terms of the Budapest Treaty on the international recognition of the deposit of microorganisms for the purposes of patent procedure.

In specific embodiments, the polynucleotides of the invention are at least 15, at least 30, at least 50, at least 100, at least 125, at least 500, or at least 1000 continuous nucleotides but are less than or equal to 300 kb, 200 kb, 100 kb, 50 kb, 15 kb, 10 kb, 7.5kb, 5 kb, 2.5 kb, 2.0 kb, or 1 kb, in length. In a further embodiment, polynucleotides of the invention comprise a portion of the coding sequences, as disclosed herein, but do not comprise all or a portion of any intron. In another embodiment, the polynucleotides comprising coding sequences do not contain coding sequences of a genomic flanking gene (i.e., 5' or 3' to the gene of interest in the genome). In other embodiments, the polynucleotides of the invention do not contain the coding sequence of more than 1000, 500, 250, 100, 50, 25, 20, 15, 10, 5, 4, 3, 2, or 1 genomic flanking gene(s).

[69] A "polynucleotide" of the present invention also includes those polynucleotides capable of hybridizing, under stringent hybridization conditions, to sequences contained in SEQ ID NO:X, or the complement thereof (e.g., the complement of any one, two, three, four, or more of the polynucleotide fragments described herein), the polynucleotide sequence delineated in columns 7 and 8 of Table 1A or the complement thereof, the

polynucleotide sequence delineated in columns 8 and 9 of Table 2 or the complement thereof, and/or cDNA sequences contained in Clone ID NO:Z (e.g., the complement of any one, two, three, four, or more of the polynucleotide fragments, or the cDNA clone within the pool of cDNA clones deposited with the ATCC, described herein), and/or the polynucleotide sequence delineated in column 6 of Table 1C or the complement thereof. "Stringent hybridization conditions" refers to an overnight incubation at 42 degree C in a solution comprising 50% formamide, 5x SSC (750 mM NaCl, 75 mM trisodium citrate), 50 mM sodium phosphate (pH 7.6), 5x Denhardt's solution, 10% dextran sulfate, and 20  $\mu$ g/ml denatured, sheared salmon sperm DNA, followed by washing the filters in 0.1x SSC at about 65 degree C.

- [70] Also contemplated are nucleic acid molecules that hybridize to the polynucleotides of the present invention at lower stringency hybridization conditions. Changes in the stringency of hybridization and signal detection are primarily accomplished through the manipulation of formamide concentration (lower percentages of formamide result in lowered stringency); salt conditions, or temperature. For example, lower stringency conditions include an overnight incubation at 37 degree C in a solution comprising 6X SSPE (20X SSPE = 3M NaCl; 0.2M NaH<sub>2</sub>PO<sub>4</sub>; 0.02M EDTA, pH 7.4), 0.5% SDS, 30% formamide, 100 ug/ml salmon sperm blocking DNA; followed by washes at 50 degree C with 1XSSPE, 0.1% SDS. In addition, to achieve even lower stringency, washes performed following stringent hybridization can be done at higher salt concentrations (e.g. 5X SSC).
- [71] Note that variations in the above conditions may be accomplished through the inclusion and/or substitution of alternate blocking reagents used to suppress background in hybridization experiments. Typical blocking reagents include Denhardt's reagent, BLOTTO, heparin, denatured salmon sperm DNA, and commercially available proprietary formulations. The inclusion of specific blocking reagents may require modification of the hybridization conditions described above, due to problems with compatibility.
- [72] Of course, a polynucleotide which hybridizes only to polyA+ sequences (such as any 3' terminal polyA+ tract of a cDNA shown in the sequence listing), or to a complementary stretch of T (or U) residues, would not be included in the definition of

"polynucleotide," since such a polynucleotide would hybridize to any nucleic acid molecule containing a poly (A) stretch or the complement thereof (e.g., practically any double-stranded cDNA clone generated using oligo dT as a primer).

The polynucleotide of the present invention can be composed of any polyribonucleotide or polydeoxribonucleotide, which may be unmodified RNA or DNA or modified RNA or DNA. For example, polynucleotides can be composed of single- and double-stranded DNA, DNA that is a mixture of single- and double-stranded regions, single- and double-stranded RNA, and RNA that is mixture of single- and double-stranded regions, hybrid molecules comprising DNA and RNA that may be single-stranded or, more typically, double-stranded or a mixture of single- and double-stranded regions. In addition, the polynucleotide can be composed of triple-stranded regions comprising RNA or DNA or both RNA and DNA. A polynucleotide may also contain one or more modified bases or DNA or RNA backbones modified for stability or for other reasons. "Modified" bases include, for example, tritylated bases and unusual bases such as inosine. A variety of modifications can be made to DNA and RNA; thus, "polynucleotide" embraces chemically, enzymatically, or metabolically modified forms.

In specific embodiments, the polynucleotides of the invention are at least 15, at least 30, at least 50, at least 100, at least 125, at least 500, or at least 1000 continuous nucleotides but are less than or equal to 300 kb, 200 kb, 100 kb, 50 kb, 15 kb, 10 kb, 7.5kb, 5 kb, 2.5 kb, 2.0 kb, or 1 kb, in length. In a further embodiment, polynucleotides of the invention comprise a portion of the coding sequences, as disclosed herein, but do not comprise all or a portion of any intron. In another embodiment, the polynucleotides comprising coding sequences do not contain coding sequences of a genomic flanking gene (i.e., 5' or 3' to the gene of interest in the genome). In other embodiments, the polynucleotides of the invention do not contain the coding sequence of more than 1000, 500, 250, 100, 50, 25, 20, 15, 10, 5, 4, 3, 2, or 1 genomic flanking gene(s).

"SEQ ID NO:X" refers to a polynucleotide sequence described in column 5 of Table 1A, while "SEQ ID NO:Y" refers to a polypeptide sequence described in column 10 of Table 1A. SEQ ID NO:X is identified by an integer specified in column 6 of Table 1A. The polypeptide sequence SEQ ID NO:Y is a translated open reading frame (ORF) encoded by polynucleotide SEQ ID NO:X. The polynucleotide sequences are shown in the

sequence listing immediately followed by all of the polypeptide sequences. Thus, a polypeptide sequence corresponding to polynucleotide sequence SEQ ID NO:2 is the first polypeptide sequence shown in the sequence listing. The second polypeptide sequence corresponds to the polynucleotide sequence shown as SEQ ID NO:3, and so on.

**[76]** The polypeptide of the present invention can be composed of amino acids joined to each other by peptide bonds or modified peptide bonds, i.e., peptide isosteres, and may contain amino acids other than the 20 gene-encoded amino acids. polypeptides may be modified by either natural processes, such as posttranslational processing, or by chemical modification techniques which are well known in the art. Such modifications are well described in basic texts and in more detailed monographs, as well Modifications can occur anywhere in a as in a voluminous research literature. polypeptide, including the peptide backbone, the amino acid side-chains and the amino or carboxyl termini. It will be appreciated that the same type of modification may be present in the same or varying degrees at several sites in a given polypeptide. Also, a given polypeptide may contain many types of modifications. Polypeptides may be branched, for example, as a result of ubiquitination, and they may be cyclic, with or without branching. Cyclic, branched, and branched cyclic polypeptides may result from posttranslation natural processes or may be made by synthetic methods. Modifications include acetylation, acylation, ADP-ribosylation, amidation, covalent attachment of flavin, covalent attachment of a heme moiety, covalent attachment of a nucleotide or nucleotide derivative, covalent attachment of a lipid or lipid derivative, covalent attachment of phosphotidylinositol, cross-linking, cyclization, disulfide bond formation, demethylation, formation of covalent cross-links, formation of cysteine, formation of pyroglutamate, formylation, gamma-carboxylation, glycosylation, GPI anchor formation, hydroxylation, iodination, methylation, myristoylation, oxidation, pegylation, proteolytic processing, phosphorylation, prenylation, racemization, selenoylation, sulfation, transfer-RNA mediated addition of amino acids to proteins such as arginylation, and ubiquitination. (See, for instance, PROTEINS - STRUCTURE AND MOLECULAR PROPERTIES, 2nd Freeman and Company, New York (1993); Ed., T. E. Creighton, W. H. POSTTRANSLATIONAL COVALENT MODIFICATION OF PROTEINS, B. Johnson, Ed., Academic Press, New York, pgs. 1-12 (1983); Seifter et al., Meth. Enzymol. 182:626-646 (1990); Rattan et al., Ann. N.Y. Acad. Sci. 663:48-62 (1992)).

"SEQ ID NO:X" refers to a polynucleotide sequence described, for example, in Tables 1A, 1B or 2, while "SEQ ID NO:Y" refers to a polypeptide sequence described in column 11 of Table 1A and or column 6 of Table 1B. SEQ ID NO:X is identified by an integer specified in column 4 of Table 1B. The polypeptide sequence SEQ ID NO:Y is a translated open reading frame (ORF) encoded by polynucleotide SEQ ID NO:X. "Clone ID NO:Z" refers to a cDNA clone described in column 2 of Table 1A and/or 1B.

[78] "A polypeptide having functional activity" refers to a polypeptide capable of displaying one or more known functional activities associated with a full-length (complete) protein. Such functional activities include, but are not limited to, biological activity, antigenicity [ability to bind (or compete with a polypeptide for binding) to an anti-polypeptide antibody], immunogenicity (ability to generate antibody which binds to a specific polypeptide of the invention), ability to form multimers with polypeptides of the invention, and ability to bind to a receptor or ligand for a polypeptide.

[79] The polypeptides of the invention can be assayed for functional activity (e.g. biological activity) using or routinely modifying assays known in the art, as well as assays described herein. Specifically, one of skill in the art may routinely assay secreted polypeptides (including fragments and variants) of the invention for activity using assays as described in the examples section below.

[80] "A polypeptide having biological activity" refers to a polypeptide exhibiting activity similar to, but not necessarily identical to, an activity of a polypeptide of the present invention, including mature forms, as measured in a particular biological assay, with or without dose dependency. In the case where dose dependency does exist, it need not be identical to that of the polypeptide, but rather substantially similar to the dose-dependence in a given activity as compared to the polypeptide of the present invention (i.e., the candidate polypeptide will exhibit greater activity or not more than about 25-fold less and, preferably, not more than about tenfold less activity, and most preferably, not more than about three-fold less activity relative to the polypeptide of the present invention).

## TABLE 1A

Last AA of ORF 141 6	$\infty$			_		
	18	25	41	139	30	20
Last AA First of AA of Sig Secreted Pep Portion 6 7	14	10	24	29	21	12
	13	6	23	28	20	11
First Last AA AA of of Sig Sig Pep Pep 1 6	<b>-</b>	-  -	1	1	1	1
AA SEQ DD NO: Y Y S15 516 516 517	518	519	520	521	522	523
of of First AA of Signal Pep 83 83	203	250	217	250	347	238
5' NT of Start Codon 1448	135	Ci	217	250	347	
3' NT of Clone Seq. 605	1493	888	3239	299	2318	330
S' NT of Clone Seq. 44	-  -	-	-	-	1	1
L	1493	888	3239	299	2318	330
SEQ SEQ 12 11 X :: 12 12 12 13 13 13 13 13 13 13 13 13 13 13 13 13			16	17	18	19
Vector Uni-ZAP XR Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pSport1	pSport1	pBluescript
ATCC Deposit No:Z and Date 203917 04/08/99 203959 04/26/99			04/29/99 203979 04/29/99	203917 04/08/99	203917 04/08/99	203979 04/29/99
cDNA Clone ID H6BSF56 H6EDM64	HACAB68	HACBJ56	HACBS22	HADDE71	HADDJ13	HADMB15
Gene No. 2	J 4	5	9	7	∞	6

5'  NT   3'  NT
of
Clone Clone
Seq.
743
1284
2890
785
874
2440
1346
1237
2345
2536
2182

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		Last	AA		ORF	1167		23		3		89		42		37		17		42		16		124		
		First	AA of		Portion	23		16				21		17		24				20		19		28		
	First Last	AA		Sig	Pep	22		15				20		16		23				19		18		27		
	First	AA	of	Sig	Pep	1		_		1				1		1		1				-		-		
	AA	SEQ	Α		Y	535		536		537		538		539		540		541		542		543		544		
5' NT	Jo	First	AA of	Signal	Pep	8		250		262		18		268		296		271		93		521		17		
		5' NT	Jo	Start	Codon			250				18		268		296		271		93		521		17		
	3' NT	of	Clone	Seq.	1	4802		602		969		1380		903		1809		934		850		1596		720		
	5' NT 3' NT	of	Clone Clone	Seq.	1	7		1		1		_		1		95		-		1		293		1		
			Total	NT	Seq.	5143		209		969		1380		903		1809		934		850		1713		720	•	
	K	SEQ	А	NO:	X	31		32		33		34		35		36		37		38		39		40		
					Vector	pSport1		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		pSport1	•	pSport1		
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				Gene	No.	21		22		23		24		25		26		27		28		29		30		

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		First	AA of	Secreted	Portion	28			70		70				6			17		25		16		25		17	
	Last	AA		Sig		27			19		19				∞			16		24		15		24		16	
	First Last	AA	of	Sig	Pep	1			1		_				<b>-</b>	_		_		<del>-</del>		<del></del>		_		-	
	AA	SEQ			Y	1015			545		546		547		548			549		550		551		552		553	
5' NT	Jo	First	AA of	Signal NO:	Pep	1033			351		671		1016		550			110		28		1877		1036		586	
		5' NT	of	Start	Codon	1033			351		671									28		1877		1036			
	3, NT	Jo	Clone	Seq.		1747			<b>687</b>		1007		1829		802			069		1647		2392		1545		619	
	5' NT 3' NT	Jo	Clone Clone	Seq.		1027			1		320		764					1		_		1612		808		Ţ	
			Total	ZZ	Seq.	2878			<i>L</i> 89		1001		1856		802			069		1647		2392		1782		619	
	N	SEQ	А	NO:	X	511			41		42		43		44			45		46		47		48		49	
					Vector	pSport1	1		pSport1		pSport1	•	Uni-ZAP XR		Uni-ZAP XR			Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR	
		ATCC	Deposit	No:Z	and Date	PTA-	794	09/27/99	203917	04/08/99	203917	04/08/99	203917	04/08/69	PTA-	793	09/27/99	203917	04/08/99	503959	04/26/99	203917	04/08/99	203917	04/08/99	203917	04/08/99
				cDNA	$\sim$	HBCJL35			HBDAB91		HBDAB91		HBGBC29		HBGNC72			HBHAA05		HBHAA81		HBIAA59		HBIAC29		HBICW51	
				Gene	No.	30			31		32		33		34			35		36		37	-	38		39	

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		Last			ORF	34	23	31		733		107	×	·	36	ľ	<u>~</u>	$\perp$	77			31		4	
		First	AA of	Secreted	Portion	28	14			2		38			25				17			19			
	Last	AA	Jo	Sig	Pep	27	13			-		37			24				16			18		_	
	First Last	AA	of	Sig	Pep		1	1		-			-	T	1	,	<del>-</del>		<del></del>			-			
	AA	SEQ	А	$\sim$		554	555	556		557		558	550	777	260	,	561		562			563		564	
5, NT	of	First	AA of	Signal	Pep	84	137	47		289		1032	121	171	115	!	430		321			2/8		170	
		5' NT	Jo	Start	Codon	84	137	47		685		1032			115							28			
	3, NT	of	Clone	Seq.		1665	892	1135		3202		2325	627	/50	750		543		637			1629		1076	
	5' NT 3' NT	Jo	Clone Clone	Seq.		1	-	-		2270		968	-	<b>⊣</b>	-		_					1		1	
	•		Total	N	Seq.	1693	1685	1135		3208		2325	637	/50	750		543		637			1629		1076	
	Z	SEQ	А	NO:	X	20	51	52		53		54	3	C	56		27		28			59		09	
					Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR	TI.: 7AD VD	UNI-CAF AK	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR			Uni-ZAP XR		pBluescript	
		ATCC	Deposit	No:Z	and Date	203917 04/08/99	203917	203917	04/08/99	203917	04/08/99	203917	200017	203917 04/08/99	203979	04/72/99	203917	04/08/99	PTA-	181	66/L0/90	203979	04/29/99	203917	04/08/99
	•			cDNA	$\overline{}$	HBJAB02	HBJAC65	HRIBM12		HBJCR46		HBJDS79	73711011	нылмэө	HBJEL16		HBJFK45		HBJIG20			HBJKD16		HBMBM96	
				Gene	No.	40	41	42	!	43		44	,	5	46		47		48			49		50	

ATCC ODNA OPPOSIT CONA NO:Z CONG ID ATCC OPPOSIT CONG ID ATCC OPPOSIT CONG ID AND Date O4/08/99 HBMTX26 O4/08/99 HBMTX48 CO3917 HBMTY48 CO3917 HBMTY48 CO3917 HBMWE61 CO4/08/99 HBMWE61 CO3917 CO1-ZAP XR CO2017 CO	Γ									5, NT	Γ				
ATCC   ATCC   ACC   AC					NT		5' NT	3, NT		of		First	Last		
CDNA         No.Z         No.Z         No.Z         No.Z         No.Z         No.Z         No.Z         No.Z         Seq. Seq. Start         Signal Signal NO: Sig			ATCC		SEQ		Jo		5' NT				AA	First	Last
CDNA         No.Z         No.Z         No.Z         No.Z         No.Z         No.Z         No.Z         Seq.         Stat.         Stat. <td></td> <td></td> <td>Deposit</td> <td></td> <td>Α</td> <td>Total</td> <td>Clone</td> <td>Clone</td> <td></td> <td>AA of</td> <td>А</td> <td></td> <td>Jo</td> <td>AA of</td> <td>AA</td>			Deposit		Α	Total	Clone	Clone		AA of	А		Jo	AA of	AA
Clone ID         and Date         Vector         X         Seq.         Codon         Pep         Y         Pep         Portion           HBMBX01         203917         pBluescript         61         1652         179         1458         363         363         565         1         18         19           HBMTX1         203917         Uni-ZAP XR         62         1639         1         1639         125         155         566         1         19         20           HBMTX26         203917         Uni-ZAP XR         63         1308         1         1801         107         567         1         46         47           HBMTY48         203917         Uni-ZAP XR         64         1891         1         1891         600         660         568         1         3         3         3           HBMUTY48         203917         Uni-ZAP XR         65         726         1         726         344         344         569         1         14         47           HBMVB51         203917         Uni-ZAP XR         65         726         1         726         349         349         579         1         1         1 <t< td=""><td>ne</td><td></td><td>No:Z</td><td></td><td>NO:</td><td></td><td>Seq.</td><td></td><td>_</td><td>Signal</td><td>SO:</td><td></td><td>Sig</td><td>Secreted</td><td>of</td></t<>	ne		No:Z		NO:		Seq.		_	Signal	SO:		Sig	Secreted	of
HBMTX101         203917 (Mi-ZAP XR)         62 (639)         179 (1458)         363 (36)         565 (1)         18 (19)         19           HBMTX11         203917 (Mi-ZAP XR)         62 (639)         1 (63	Ö.	Clone ID	and Date	Vector	X	Seq.	1		Codon	Pep	Y				ORF
HBMTM11 203917 Uni-ZAP XR 62 1639 1 1639 125 125 566 1 19 20  HBMTX26 203917 Uni-ZAP XR 63 1308 1 1891 660 660 568 1 36 37  HBMTX48 203917 Uni-ZAP XR 64 1891 1 1891 660 660 568 1 36 37  HBMUH74 PTA- Uni-ZAP XR 65 118 1 118 238 238 570 1 13 14  HBMWE01 203917 Uni-ZAP XR 66 1118 1 1118 238 238 570 1 29 30  HBNAX40 203917 Uni-ZAP XR 68 1974 1469 1974 1603 573 1 29 30  HBQAB79 203917 Lambda ZAP XR 68 1331 1 1331 190 190 573 1 29 30  HBQACS7 203917 Lambda ZAP XR 67 2111 1 1311 146 146 574 1 1 18 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1		HBMBX01	203917	pBluescript	61	1652		1458	363	363	595	-	18	19	78
HBMTM11         203917         Uni-ZAP XR         62         1639         1         1639         125         125         566         1         19         20           HBMTX26         203917         Uni-ZAP XR         63         1308         1         1308         107         107         567         1         46         47           HBMTY48         203917         Uni-ZAP XR         64         1891         1         1891         660         660         568         1         36         37           HBMUH74         PTA-         Uni-ZAP XR         65         726         1         726         344         344         569         1         13         14           HBMUH74         PTA-         Uni-ZAP XR         66         1118         1         118         238         238         570         1         13         14           HBMWE61         203917         Uni-ZAP XR         67         1793         2455         2793         2497         2497         571         1         18         19           HBNB176         203917         Lambda ZAP XR         68         1974         1469         1974         160         573         1			04/08/69												
HBMTX26         203917         Uni-ZAP XR         63         1308         1         1308         107         107         567         1         46         47           HBMTY48         203917         Uni-ZAP XR         64         1891         1         1891         660         660         568         1         36         37           HBMUH74         PTA-         Uni-ZAP XR         65         726         1         726         344         344         569         1         13         14           HBMWE61         203917         Uni-ZAP XR         66         1118         1         1118         238         570         1         18         19           HBNAX40         203917         Uni-ZAP XR         67         1793         2455         2793         2497         5497         571         1         18         19           HBNBJ76         203917         Uni-ZAP XR         68         1974         1469         1974         1603         572         1         29         30           HBNBJ76         203917         Lambda ZAP         69         1331         1         131         146         1974         160         190         573	52	HBMTM11	203917	Uni-ZAP XR	62	1639	П	1639	125	125	999	<del></del>	19	70	31
HBMTX26         203917         Uni-ZAP XR         63         1308         1         1308         107         567         1         46         47           HBMTY48         203917         Uni-ZAP XR         64         1891         1         1891         660         660         568         1         36         37           HBMUH74         PTA-         Uni-ZAP XR         65         726         1         726         344         344         569         1         13         14           HBMWE61         203917         Uni-ZAP XR         66         1118         1         1118         238         238         570         1         18           HBNAX40         203917         Uni-ZAP XR         67         1745         1469         1974         1469         1974         1469         1974         1603         572         1         29         30           HBQAB79         203917         Uni-ZAP XR         68         1974         1469         1974         1603         573         1         29         30           HBQAB79         203917         Lambda ZAP         69         1331         1         1331         190         190         573 <td></td> <td></td> <td>04/08/66</td> <td></td>			04/08/66												
HBMTY48   203917   Uni-ZAP XR   64   1891   1   1891   660   660   568   1   36   37     HBMUH74   PTA-	53	HBMTX26	203917	Uni-ZAP XR	63	1308	-	1308	107	107	267	_	46	47	68
HBMUH748   203917   Uni-ZAP XR   64   1891   1   1891   660   568   1   36   37     HBMUH74   PTA-			04/08/99												
HBMUH74 PTA- Uni-ZAP XR 65 726 1 726 344 344 569 1 13 14  HBMWE61 203917 Uni-ZAP XR 66 1118 1 1118 238 238 570 1 18 19  HBNAX40 203917 Uni-ZAP XR 68 1974 1469 1974 1603 572 1 29 30  HBOAB79 203917 Lambda ZAP 69 1331 1 1331 190 190 573 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	4	HBMTY48	203917	Uni-ZAP XR	64	1891		1891	099	099	268	-	36	37	45
HBMUH74         PTA-         Uni-ZAP XR         65         726         1         726         344         344         569         1         13         14           181         181         181         26/07/99         26/07/99         26/07/99         26/07/99         2793         2452         2793         2497         2497         570         1         18         19           HBNAX40         203917         Uni-ZAP XR         68         1974         1469         1974         1603         572         1         29         30           HBNBJ76         203917         Uni-ZAP XR         68         1974         1469         1974         1603         572         1         29         30           HBQAB79         203917         Lambda ZAP         69         1331         1         1331         190         190         573         1         29         30           HBQAC57         203917         Lambda ZAP         70         2111         1         2111         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146 <td></td> <td></td> <td>04/08/99</td> <td></td>			04/08/99												
181       06/07/99       1118       1118       238       238       570       1       1         203917       Uni-ZAP XR       66       1118       1       1118       238       570       1       18       19         203917       Uni-ZAP XR       68       1974       1469       1974       1603       572       1       29       30         203917       Lambda ZAP       69       1331       1       1331       190       190       573       1       29       30         203917       Lambda ZAP       70       2111       1       2111       146       146       174       16       374       1       29       30         203917       Lambda ZAP       70       2111       1       2111       146       146       574       1       29       30         203917       Lambda ZAP       70       2111       1       2111       146 <td< td=""><td>55</td><td>HBMUH74</td><td>PTA-</td><td>Uni-ZAP XR</td><td>65</td><td>726</td><td></td><td>726</td><td>344</td><td>344</td><td>569</td><td><u></u></td><td>13</td><td>14</td><td>78</td></td<>	55	HBMUH74	PTA-	Uni-ZAP XR	65	726		726	344	344	569	<u></u>	13	14	78
HBMWE61       203917       Uni-ZAP XR       66       1118       1       1118       238       238       570       1         04/08/99       04/08/99       104/08/99       203917       Uni-ZAP XR       67       2793       2455       2793       2497       2497       571       1       18       19         HBNBJ76       203917       Uni-ZAP XR       68       1974       1469       1974       1603       572       1       29       30         HBQAB79       203917       Lambda ZAP       69       1331       1       1331       190       190       573       1       29       30         HBQAC57       203917       Lambda ZAP       70       2111       1       2111       146       146       174       146       174       146 </td <td></td> <td></td> <td>181</td> <td></td>			181												
HBMWE61         203917         Uni-ZAP XR         66         1118         1         1118         238         570         1           04/08/99         HBNAX40         203917         Uni-ZAP XR         67         2793         2455         2793         2497         2497         571         1         18         19           HBNBJ76         203917         Uni-ZAP XR         68         1974         1469         1974         1603         572         1         29         30           HBQAB79         203917         Lambda ZAP         69         1331         1         1331         190         190         573         1         29         30           HBQAC57         203917         Lambda ZAP         70         2111         1         2111         146         146         574         1         2           04/08/99         II         203917         Lambda ZAP         70         2111         1         2111         146         146         574         1         2         1			66/L0/90												
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HBNAX40 203917 Uni-ZAP XR 68 1974 1469 1974 1603 572 1 29 30 4/08/99 HBQAB79 203917 Lambda ZAP 69 1331 1 1211 146 146 574 1 6			04/08/99		į		0466	2010	2000	7070	57.1	-	101	Q.	ę
HBNBJ76 203917 Uni-ZAP XR 68 1974 1469 1974 1603 572 1 29 30 04/08/99		HBNAX40	203917	Uni-ZAP XR	/0		7422	56/7	7447	7447	1/0	_	01	<del>.</del>	<del>,</del>
HBNBJ76         203917         Uni-ZAP XR         68         1974         1469         1974         1603         572         1         29         30           04/08/99         104/08/99         II         1331         1         1331         190         190         573         1         1           HBQAC57         203917         Lambda ZAP         70         2111         1         2111         146         146         574         1           04/08/99         II         2111         1         2111         146         146         574         1			04/08/99												
HBQAB79       203917       Lambda ZAP L	<u>∞</u>	HBNBJ76	203917	Uni-ZAP XR	89			1974		1603	572	-	59	30	89
HBQAB79 203917 Lambda ZAP 69 1331 1 1331 190 190 573 1			04/08/99												
04/08/99       II       1       2111       1       446       574       1         HBQAC57       203917       Lambda ZAP       70       2111       1       2111       146       574       1	6	HBQAB79	203917	Lambda ZAP		1331	1	1331	190	190	573	_			Ξ
HBQAC57 203917 Lambda ZAP 70 2111 1 2111 146 146 574 1 04/08/99 II		,	04/08/99	П											
	0	HBQAC57	203917	Lambda ZAP	_	2111	_	2111	146	146	574				53
			04/08/99	п											

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		Last	AA	of	ORF	48			16		-		35		8		254		23		32		32		55		55	
		First	AA of	Sig Secreted	Portion	28							18		13		18		16		14		6		27		16	
	First Last	AA	Jo			27							17		12		17		15		13		8		26		15	
		AA	of	Sig	Pep	1			_				1		-		-		-		1	-	1		1		-	
	AA	SEQ	Ω	ÖN:	Y	575			576		577		578		579		580	•	581		582		583		584		585	
5' NT	Jo	First	AA of	Signal	Pep	447			119		1148		098		333		77		29	•	1588		999		177		131	
		5' NT	of	Start	Codon	447			119						333		11								177		131	
	3' NT	of	Clone Clone	Seq.		592			1010		1219		1392		813		1896		1276		1806		1732		1419		1052	
	5' NT 3' NT	of		Seq.		129			41		-		628		1		1		1		1347		282		-		1	
			Total	Z	Seq.	592			1010		1219		1392		813		1896		1276		1807		1732		1419		1052	
	N	SEQ	П	NO:	X	71			72		73		74		75		9/		77		78		79		08		81	
					Vector	Uni-ZAP XR			ZAP Express		ZAP Express		Uni-ZAP XR															
	•	ATCC	Deposit	No:Z	and Date	-VLA	181	66/10/90	203917	04/08/99	203917	04/00/99	203917	04/08/99	203917	04/08/99	203917	04/08/99	203979	04/29/99	203917	04/08/99	203917	04/08/99	203917	04/08/99	_	04/08/99
				cDNA	Clone ID	HBSAK32			HBXCM66		HBXCX15	Т	HCDCY76		HCDDL48		HCE1G78		HCE2H52		HCE3B04		HCE5F78		HCEDR26		HCEEE79	
				Gene	No.	61			62		63		49		65		99		<i>L</i> 9		89		69		70		71	

		Last	A	بيه	ORF	3	43		265		15		2				7:		8		14		<b>—</b>			_	
		<u> </u>	AA	d of		23	4		7				25				147		∞		1		31			47	
		First	AA of	Secreted	Portion	91	31		17				17				24		32				20			26	
	Last	AA		Sig		15	30		16				16				23		31				19			25	
	First	AA	of	Sig	Pep	1	-		-		-		-		-		_		ij		-		Ī			1	
	AA	SEQ	А	Ö.	Υ	985	587		588		589		590		591		592		593		594		595			596	
5' NT	Jo	First	AA of	Signal NO:	Pep	111	209		215		237		101		384		304		539		101		1145		,	31	
		5' NT	Jo	Start	Codon		209		215		237		101			-					101		1145			31	
	3' NT	Jo	Clone	Seq.		992	1229		1781		1305		1434		735		1359		2190		746		1633			1796	
	5' NT 3' NT	of	Clone Clone	Seq.		1	-		4		1		1		1		62		334		1		1031			9//	
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	NT	SEQ	О	ö	X	82	83		84		85		98		87		88		68		06		91			92	
					Vector	Uni-ZAP XR	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		pSport1		pSport1		pSport1		Uni-ZAP XR		Lambda ZAP	П	pBluescript			Lambda ZAP	П
		ATCC	Deposit	No:Z	and Date	203917 04/08/99	203917	04/08/66	203917	04/08/66	203917	04/08/66	203917	04/08/99	203917	04/08/69	203979	04/29/99	203917	04/08/99	203917	04/08/99	PTA-	181	66/L0/90	203917	04/08/99
				cDNA	Clone ID	нсее025	HCEEU18		HCEFZ82		HCEGX05		HCFLN88		HCFLT90		HCHAB84		HCMSX51		HCNC011		HCNSD29			нсовн72	
				Gene	No.	72	73		74		75		9/		77		78		79		80		81			82	

AA First Last SEQ AA AA First ID of of AA of NO: Sig Sig Secreted Y Pep Pep Portion 597 1 20 21 599 1										5' NT					
ATCC         SEQ         of         of         of         S' NT         First         SEQ         AA of         DD         of         AA of         AA of         DD         of         AA					Z		5' NT	3' NT		of	AA	First	Last		
Deposit         D         Total         Clone         Clone         of         AA of         D         of         AA of           No.2         No.2         NT         Seq.         Seq.         Stat         Signal         NO.         Sig         Sig         Secreted           203979         Lambda ZAP         93         2166         632         1455         782         597         1         20         21           203979         Lambda ZAP         94         1287         1         1287         782         598         1         20         21           203979         Lambda ZAP         94         1287         1         1287         728         598         1         20         21           203979         Lambda ZAP         95         1929         606         1929         815         815         1         20         21           203971         Lambda ZAP XR         96         788         1         788         141         600         1         36         37           203917         Lambda ZAP XR         97         1264         101         1142         191         191         601         1         23         24			ATCC		SEQ		Jo		5'NT	First	SEQ	AA	AA	First	Last
No:Z         NO: NT Seq.         Seq. Seq. Seq. Start         Signal Operation of Date (Codon)         NO: No. NT Seq. Seq. Seq. Seq. Seq. Seq. Seq. Seq.			Deposit		А	Total	Clone	Clone	Jo	AA of		of	of	AA of	AA
and Date         Vector         X         Seq.         A         Codon         Pep         Y         Pep         Pep         Potion           203979         Lambda ZAP         93         2166         632         1455         782         782         597         1         20         21           203917         Lambda ZAP         94         1287         1         1287         728         598         1         20         21           203979         Lambda ZAP         95         1929         606         1929         815         815         599         1         20         21           203917         Uni-ZAP XR         96         788         1         788         1         141         600         1         36         37           203917         Uni-ZAP XR         96         788         1         788         1         141         600         1         36         37           203917         Uni-ZAP XR         96         788         1         788         363         363         363         363         363         363         363         363         363         363         363         363         363         363<		cDNA	No:Z		SON:	N		Seq.	Start	Signal	NO:			Secreted	Jo
203979         Lambda ZAP         93         2166         632         1455         782         782         597         1         20         21           203917         Lambda ZAP         94         1287         1         1287         728         598         1         2         21           203979         Lambda ZAP         95         1929         606         1929         815         815         599         1         8         7           203979         Lambda ZAP         96         788         1         788         141         600         1         36         37           203917         Uni-ZAP XR         96         788         1         188         141         600         1         36         37           203917         Uni-ZAP XR         96         788         1         188         191         601         1         1         2           203917         Uni-ZAP XR         97         1264         101         1142         191         191         601         1         2         2           203917         AP Express         10         530         1         597         230         230         1		Clone ID	and Date		X	Seq.			Codon		Y	Pep	Pep		ORF
04/29/99         II         PA         1287         1         1287         728         598         1           203979         I ambda ZAP         94         1287         1         1287         728         598         1           203979         I ambda ZAP         95         1929         606         1929         815         815         599         1           203917         Uni-ZAP XR         96         788         1         788         141         600         1         36         37           203917         Uni-ZAP XR         96         788         1         788         141         600         1         36         37           203917         Uni-ZAP XR         97         1264         101         1142         191         191         601         1         2         23           203917         PSport1         98         892         1         892         363         602         1         22         23           203917         ZAP Express         100         530         1         597         230         230         1         23         24           04/08/99         20395         204         1		<b>9622021</b>	203979	Lambda ZAP	93	2166	632	1455	782	782	597	1	20	21	45
203917         Lambda ZAP         94         1287         1         1287         728         598         1           204/08/99         II         203979         Lambda ZAP         95         1929         606         1929         815         815         599         1         7           203979         Lambda ZAP XR         96         788         1         788         141         600         1         36         37           203917         Uni-ZAP XR         97         1264         101         1142         191         601         1         1         2         23           203917         Uni-ZAP XR         97         1264         101         1142         191         601         1         1         2           203917         Apport1         98         892         1         892         363         363         603         1         2         23           203917         ZAP Express         100         530         1         597         230         189         604         1         18         19           203957         ZAP Express         101         1143         578         1136         598         596			04/29/99	П											<u> </u>
04/08/99         Lambda ZAP XR         95         1929         606         1929         815         815         599         1           203979         Lambda ZAP XR         96         788         1         788         141         600         1         36         37           203917         Uni-ZAP XR         96         788         1         188         1141         600         1         36         37           203917         Uni-ZAP XR         97         1264         101         1142         191         191         601         1         2           203917         Uni-ZAP XR         97         1264         101         1142         191         191         601         1         2           203917         ZAP Express         99         597         1         892         363         363         602         1         22         23           203917         ZAP Express         100         530         1         530         189         604         1         18         19           203957         ZAP Express         101         1143         578         1136         598         605         1         30         31 <td>_</td> <td>нсосл56</td> <td>203917</td> <td>Lambda ZAP</td> <td>94</td> <td>1287</td> <td>1</td> <td>1287</td> <td></td> <td>728</td> <td>869</td> <td></td> <td></td> <td></td> <td>1</td>	_	нсосл56	203917	Lambda ZAP	94	1287	1	1287		728	869				1
203979         Lambda ZAP         95         1929         606         1929         815         899         1           203917         Uni-ZAP XR         96         788         1         788         141         600         1         36         37           203917         Uni-ZAP XR         97         1264         101         1142         191         601         1         1         2           203917         Uni-ZAP XR         97         1264         101         1142         191         601         1         1         2           203917         pSport1         98         892         1         892         363         363         602         1         2         23           04/08/99         203917         ZAP Express         100         530         1         597         230         230         603         1         23         24           04/08/99         203957         ZAP Express         101         1143         578         1136         598         605         1         30         31           203917         ZAP Express         102         402         150         389         256         256         606 <td>- 1</td> <td></td> <td>04/08/99</td> <td></td> <td></td> <td></td> <td>_</td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td>	- 1		04/08/99				_								
203917         Uni-ZAP XR         96         788         1         788         1         414         600         1         36         37           04/08/99         203917         Uni-ZAP XR         97         1264         101         1142         191         191         601         1         2           203917         Do4/08/99         892         1         892         363         363         602         1         2         23           203917         ZAP Express         99         597         1         597         230         230         603         1         23         24           203917         ZAP Express         100         530         1         530         189         604         1         18         19           203957         ZAP Express         101         1143         578         1136         598         605         1         30         31           203917         ZAP Express         102         402         150         389         256         256         606         1         35         36           203917         ZAP Express         103         471         1         410         607	4	ICQCM24	203979 04/29/99		95	1929		1929	815	815	599	-			38
04/08/99       Amount of the color of the c	_	1	203917		96	788	-	788		141	009	1	36	37	145
203917         Uni-ZAP XR         97         1264         101         1142         191         601         1         1         2           04/08/99         Sport1         98         892         1         892         363         363         602         1         22         23           04/08/99         203917         ZAP Express         99         597         1         597         230         230         603         1         23         24           04/08/99         Alional Salaria         1143         578         1136         598         604         1         18         19           04/08/99         Alional Salaria         1143         578         1136         598         598         605         1         30         31           203917         AP Express         102         402         150         389         256         256         606         1         35         36           203917         AP Express         103         471         1         410         607         1         35         36			04/08/69												
203917         pSport1         98         892         1         892         363         363         602         1         22         23           203917         ZAP Express         99         597         1         597         230         230         603         1         23         24           203917         ZAP Express         100         530         1         530         189         604         1         18         19           203957         ZAP Express         101         1143         578         1136         598         598         605         1         30         31           203957         ZAP Express         102         402         150         389         256         256         606         1         35         36           203917         ZAP Express         103         471         1         471         410         607         1         35         36	_	HCRBF72	203917	Uni-ZAP XR	62	1264	101	1142	191	191	601	1	1	2	211
04/08/99       203917       ZAP Express       99       597       1       597       230       603       1       23       24         203917       ZAP Express       100       530       1       530       189       604       1       18       19         203917       ZAP Express       101       1143       578       1136       598       598       605       1       30       31         203917       ZAP Express       102       402       150       389       256       256       606       1       35       36         203917       ZAP Express       103       471       1       471       410       607       1       35       36		HCRNF78	203917	pSport1	86	892	-	892	363	363	602	1	22	23	46
203917         ZAP Express         99         597         1         597         230         230         603         1         23         24           203917         ZAP Express         100         530         1         530         189         604         1         18         19           203957         ZAP Express         101         1143         578         1136         598         598         605         1         30         31           203917         ZAP Express         102         402         150         389         256         256         606         1         35         36           203917         ZAP Express         103         471         1         471         410         607         1         35         36			04/08/66												
04/08/99       203917       ZAP Express       100       530       1       530       189       189       604       1       18       19         04/08/99       203957       ZAP Express       101       1143       578       1136       598       598       605       1       30       31         203917       ZAP Express       102       402       150       389       256       256       606       1       35       36         203917       ZAP Express       103       471       1       471       410       607       1       35       36		HCUAF85	203917	ZAP Express	66	597	-	597	230	230	603	1	23	24	122
203917       ZAP Express       100       530       1       530       189       189       604       1       18       19         204/08/99       203957       ZAP Express       101       1143       578       1136       598       598       605       1       30       31         203917       ZAP Express       102       402       150       389       256       256       606       1       35       36         203917       ZAP Express       103       471       1       471       410       607       1       35       36			04/08/99												
04/08/99       203957       ZAP Express       101       1143       578       1136       598       605       1       30       31         04/26/99       203917       ZAP Express       102       402       150       389       256       256       606       1       35       36         203917       ZAP Express       103       471       1       471       410       607       1       7		HCUCF89	203917	ZAP Express	100	530	-	530	189	189	604	1	18	19	59
203957       ZAP Express       101       1143       578       1136       598       598       605       1       30       31         04/26/99       203917       ZAP Express       102       402       150       389       256       256       606       1       35       36         203917       ZAP Express       103       471       1       471       410       607       1       7			04/08/99												
04/26/99       203917       ZAP Express       102       402       150       389       256       256       606       1       35       36         04/08/99       203917       ZAP Express       103       471       1       471       410       607       1	ш,	4CUCK44	203957	ZAP Express	101	1143	578	1136	869	598	909	-	30	31	99
203917 ZAP Express 102 402 150 389 256 256 606 1 35 36 36 04/08/99 471 1 471 471 410 607 1	- 1		04/26/99												
04/08/99       203917     ZAP Express     103     471     1     471     410     607     1       04/08/99	_	HCUDD64	203917	ZAP Express	102	402	150	389	256	256	909	-	35	36	49
203917 ZAP Express 103 471 1 471 410 607 1 04/08/99			04/08/69												
04/08/99	ш	ICWAE64	203917	ZAP Express	103	471	_	471		410	209	1			S
	ı		04/08/99										-		

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		Last	AA		ORF	22		1		56		12		55		120			70		425			52		7	
		First	AA of		Portion	10				56		12		39		18					70			53			
	Last	AA	of	Sig	Pep	6				25		Ξ		38		17					19			28			
	First Last	AA	Jo	Sig	Pep	1		-		-		_							-		_			1			
	AA	SEQ		NO:	Y	809		609		610		611		612		613		:	614		615			919		617	
5' NT	Jo	First SEQ	AA of	Signal	Pep	282		333		48		416		172		23			279		186			220		161	
		5' NT	Jo	Start	Codon	282		333		48				172		23			279		186			220		191	
	3' NT	of	Clone	Seq.		467		761		943		497		1536		1550			1997		2882			1889		2187	
	5' NT 3' NT	of	Clone Clone	Seq.		1		3				1		1		1			1		3			1		1	
			Total	NT	Seq.	467		761		943		497		1536		1550			1661		2882			1904		113 2187	
	IN	SEQ	О	ÖN:	X	104		105		106		107		108		109			110		111			112		113	
					Vector	ZAP Express	_	ZAP Express		pCMVSport	2.0	pCMVSport	2.0	pCMVSport	3.0	pCMVSport	3.0		pCMVSport	3.0	pCMVSport	3.0		pCMVSport	3.0	pCMVSport	3.0
		ATCC	Deposit	No:Z	and Date	203917	04/08/99	203917	04/08/99	203917	04/08/99	203917	04/08/99	203960	04/26/99	PTA-	794	09/27/99	203960	04/26/99	PTA-	181	66/L0/90	203960	04/26/99	203918	04/08/99
				cDNA	$\sim$	HCWFU39		HCWUL09		HDHAA42		HDHEB76		HDPCW16		HDPDI72			HDPDJ58		HDPFF10			HDPFU43		HDPFY18	
				Gene	No.	94		95		96		97		86		66			100		101			102		103	

									5' NT					
				N		5' NT 3' NT	3, NT		of	AA	AA First Last	Last		•
		ATCC		SEQ		Jo		5' NT	First	SEQ	AA	AA	First	Last
		Deposit		Ω	Total	Clone Clone	Clone	ot	AA of	А	Jo	of	AA of	AA
Gene	cDNA	No:Z		NO:	N	Seq.	Seq.		Signal	ON	Sig	Sig		Jo
No.	Clone ID	and Date	Vector	X	Seq.			Codon	Pep	Y	Pep	Pep	등	ORF
104	HDPGE24	203960	pCMVSport	114	2625	1	2625	173	173	618	_	11	12	73
		04/26/99	3.0											
105	HDPIU94	203960	pCMVSport	115	2196	21	2196	208	208	619		21	22	23
		04/26/99	3.0											1
106	HDPOC24	203960	pCMVSport	116	1777	302	1725	418	418	620		23	24	133
		04/26/99	3.0											
107	HDPOL37	203960	pCMVSport	117	1489	П	1489	189	189	621		32	33	62
		04/26/99	3.0											
108	HDP0076	203960	pCMVSport	118	645	Τ	645		109	622	-	15	16	16
		04/26/99	3.0											
109	HDPPD93	203960	pCMVSport	119	701	_	701	78	78	623	-			12
		04/26/99	3.0											
110	HDPPQ30	203960	pCMVSport	120	1063		1063	220	220	624	_	22	23	38
		04/26/99	3.0											1
1111	HDPPW82	203959	pCMVSport	121	552	-	552	395	395	625	-			53
		04/26/99	3.0											1
112	HDPXN20	203960	pCMVSport	122	1756	_	1756	19	61	626		70	21	41
		04/26/99	3.0											
113	нронм36	PTA-	pCMVSport	123	1547	_	1547	129	129	627	_	18	19	48
		181	3.0											
		66/1/0/90												
114	HDTAU35	203960	pCIV	124	377	<b>-</b>	377	260	260	628	-	12	13	17
		04/26/99	2.0											

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		Last	AA	of	ORF	33		20		29		9		7		7		52		13		39		42		41	_
	_	First	AA of	Sig Secreted	Portion	23		17		18		19						6				17		28		19	
	Last	AA	of	Sig	Pep	22		16		17		18						∞				16		27		18	
	First Last	AA	of	Sig	Pep					_				-				-		<del></del>		-		_		_	
	AA	SEQ	Д		Υ	629		630		631		632		633		634		635		929		637		638		639	
5, NT	Jo	First	AA of	Signal NO:	Pep	191		164		375		345		360		1731		321		10		273		770		41	
		5' NT	Jo	Start	Codon	191		164				345		360		1731		321				273		0//		41	
	3, NT		Clone	Seq.		099		829		2261		525		1663		3034		809		995		1569		1323		845	
	5' NT 3' NT	Jo	Clone Clone	Seq.		1		1				1		308		1679		-				236		638		1	
			Total	N	Seq.	099		8/9		2261		525		1663		3034		608		995		6951		1323		845	
	Z	SEO	А	NO:	X	125		126		127		128		129		130		131		132		133		134		135	
					Vector	pCMVSport	2.0	pCMVSport	2.0	pCMVSport	2.0	pCMVSport	2.0	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR	
		ATCC	Deposit	No:Z	and Date	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99
				cDNA	$\overline{}$	4		HDTFX18		HDTGW48		HDTLM18		HE2CA60		HE2CA60		HE2CH58		HE2CM39		HE2HC60		HE2P093		HE6AU52	
				Gene	No.	115		116		117		118		119		120		121		122		123		124		125	

<del>(*</del>	Last	AA	Jo	ORF	62		25		46		526		33		47		47		43		43		471		43	
	First L		Secreted	Portion O	11	+	21	$\dashv$	15	$\dashv$	27	+	61	+	34	1	34	1	76	$\dashv$			7	1	792	
Last	AA			Pep	10		70		14		56		18		33		33		25		25		_		25	
First Last	AA	of	Sig	Pep	_		_		-		_						-		_				<u> </u>		-	
AA	SEQ	А	: SQ	>	640		641		642		643		644		645		646		647		648		649		650	
5' NT of	First	AA of	Signal	Pep	295		38		171		145		210		155		155		157		1074		7		2268	
	5' NT	of	Start	Codon			38		171		145		210		155		155		157		1074				2268	
3, NT	Jo	Clone	Seq.		1526		941		298		1994		1526		1887		1887		1978		2891		4890		4085	
5' NT 3' NT	Jo	Clone Clone	Seq.	_	-		-								_		1		-		918		2918		2114	
		Total	Z	Seq.	1526		941		298		2000		1526		1887		1887		1995		2908		4907		4102	
Į	SEQ	Α	NO:	X	136		137		138		139		140		141		142		143		144		145		146	
				Vector	Uni-ZAP XR																					
	ATCC	Deposit	No:Z	and Date	203960	04/26/99	203960	04/26/99	203979	04/29/99	203979	04/29/99	203960	04/26/99	203979	04/29/99	203979	04/29/99	203979	04/29/99	203979	04/29/99	203979	04/29/99	203979	04/29/99
			cDNA	$\overline{}$	HE6CS65		HE6D092		HE6EY13		HE6FU11		HE6FV29		HE8FC45		HE8FC45		HE8FD92		HE8FD92		HE8FD92		HE8FD92	
			Gene	No.	126		127		128		129		130	`	131		132		133		134		135		136	

									5' NT					
				Z		5' NT 3' NT	3, NT		of	AA	First	Last		
		ATCC		SEQ		of	of	5' NT	First	SEQ	AA	AA	First	Last
		Deposit		А	Total	Clone Clone	Clone	Jo	AA of	А	Jo	of	AA of	AA
Gene	cDNA	No:Z		NO:	N	Seq.	Seq.	Start	Signal	NO:	Sig	Sig	Secreted	of
No.	Clone ID	and Date	Vector	X	Seq.			Codon	Pep	Y	Pep	Pep	Portion	ORF
137 E	HE8FD92	203979	Uni-ZAP XR	147	262	9861	3960	2141	2141	651	1	25	26	43
		04/29/99												
138 F	HE8SG96	PTA-	Uni-ZAP XR	148	2036	1	2036	118	118	652	1	17	18	24
		181												
		66/20/90												
139 H	HE8TY46	PTA-	Uni-ZAP XR	149	2204	1400	2204	1413	1413	653	1	18	19	187
		1838												
		00/60/50												
140 H	HE9CY05	203960	Uni-ZAP XR	150	1047	47	1047	55	55	654	-	21	22	235
		04/26/99												
141 H	HE9EA10	096807	Uni-ZAP XR	151	2114	1	2111		212	655	_	28	29	78
		04/26/99												
142 H	HE9GG20	203960	Uni-ZAP XR	152	9/9	1	9/9	319	319	959	П			6
		04/26/99												
143 F	HEBCI18	203960	Uni-ZAP XR	153	1121	713	1050	855	855	657	_	43	4	69
_		04/26/99												
144 H	HEBCY54	203960	Uni-ZAP XR	154	1189	1	1189	172	172	658		24	25	118
		04/26/99												
145 H	HEBDF77	203960	Uni-ZAP XR	155	1820	1	1820	681	681	629	1	59	30	36
		04/26/99										·		
146   H	HEBDQ91	203960	Uni-ZAP XR	156	1573	1007	1573		1211	099	1	56	30	41
		04/26/99												

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L		Last	AA	of	ORF	29		42		34		20		17		22		22		10		51			15		
		First	AA of		Portion	27		26		31						17		17				41			14		
	Last	AA	of	Sig	Pep	26		25		30						16		16				40		•	13	-	
	First Last	AA	of	Sig	Pep	1	·	1		-		-		-		1		-1		1		-			1		
	AA	SEQ	П	NO:	Y	661		662		663		664		599		999		299	·	899		699			0/9		
2, NT	Jo	First SEQ	AA of	Signal NO:	Pep	200		106		59		215		82		147		147		440		154			664		
		5' NT	Jo	Start	Codon	200		106		59		215		82		147		147		440							
	3' NT	of	Clone	Seq.		1304		1867		1125		2168	-	1260		1109		1109		1614		939			746		
	5' NT 3' NT	Jo	Clone Clone	Seq.	•	1		1		1		-		1		12		12		204		_			_		
			Total	N	Seq.	1304		1867		1125		2168		1260		1109		1109		1614		939			746		
	NT	SEQ	А	NO:	X	157		158		159		160		161		162		163		164		165		•	166		
					Vector	Uni-ZAP XR	i	Uni-ZAP XR		pSport1			Uni-ZAP XR														
		ATCC	Deposit	No:Z	and Date	203979	04/29/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203979	04/29/99	PTA-	181	66/L0/90	PTA-	181	66/1/0/90
				cDNA	Clone ID	HEBFR46		HEBGE07		HEGAU15		HELAT35		HELBU54		HELGG84		HELGG84		HEMEY47		HEOMC46			HEPBA14		
				Gene	No.	147		148		149		150		151		152		153		154		155			156		

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		Last	AA		ORF	32		20	Ş	}		22		99		7		42		15		18		9		31
		First	AA of	Secreted	Portion	27		19	96	07		17		14				18						25		23
	Last	AA	of	Sig	Pep	26		18	5	17		16		13				17						24		22
	First Last	AA	Jo	Sig	Pep	1		1	1	T		_		1		I				_		_				-
	AA	SEQ	А	NO:	Y	671		672	(4)	0/2		674		<i>5L</i> 9		9/9		<i>LL</i> 9		8/9		629		089		681
5' NT	of	First SEQ	AA of	Signal NO:	Pep	120		306	200	/27		541		767		121		136		170		216		154		40
		5' NT	of	Start	Codon	150		306	000	/67		541								170		216		154		40
	3, NT	Jo	Clone	Seq.	I	1647		829	2000	C077		1533		1778		871		887		1437		1205		1153		998
	5' NT 3' NT	of	Total Clone Clone	Seq.		1		1	Ę	C/		328		1		1		T		1		1				П
			Total	N	Seq.	1647		829	2000	C077		1533		1778		871		288		1437		1205		1153		998
	Z	SEQ		NO:	X	167		168	9	109		170		171		172		173		174		175		176		177
					Vector	pCMVSport	3.0	pCMVSport		UNI-ZAF AK		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR
		ATCC	Deposit	No:Z	and Date	203960	04/26/99	203960	04/70/42	6/6507	04/77/99	203979	04/29/99	203960	04/26/99	203979	04/29/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960
:				cDNA	$\overline{}$	неолн80		HEQBF89	T	HEICHO		HETDW58		HETEY67		HFCDW95		HFCEI04		HFCFD04		HFCFE20		HFEAY59		HFGAJ16
				Gene	No.	157		158		661		160		161		162		163		164		165		166		167

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		Last	AA	o	ORF	51	- 1	87	;	43		42	,	55	_	<u>+</u>	1	35	_[`	2		45	_	- 30 	1	<u> </u>		4
		First	AA of	Secreted	Portion	22		78		23		18		17		25	,	56		24	,	78		19				
	Last	AA		_	Pep	21	ļ	27		22		17		16		24	1	28		23	!	27		18				
	First Last	AA	of	Sig	Pep	_		_				-		1				-		_		<del></del>		_		<u> </u>		
	AA	SEQ	А	S S	×	682		683		684		685		989	ļ	<b>687</b>		889		689		069		691		692		
S' NT	of	یب	AA of	Signal	Pep	700		175		283		243		9		929		414	_	2546		203		577		<i>L</i> 9		
		5' NT	Jo	Start	Codon	700		175		283		243		9		9/9		414		2546		203		577		29		
	3, NT		Clone	Seq.		1165		1275		1157		1885		1031		2735		2644		3114		1419		1941		820		1
	5'  NT   3'  NT	Jo	Clone Clone	Seq.	_	454		110						1		341		<del></del>		2302				322		1		
			Total	N	Seq.	1280		1275		1157		1885		1031		2735		2644		3115		1419		1941		820		
	Ž	SEO	А	ON	×	178		179		180		181		182		183		184		185		186		187		188		
					Vector	pSport1		pSport1		pSport1		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		
		ATCC	Deposit	No:Z	and Date	203960	04/26/99	203960	04/26/99	203979	04/29/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	PTA-	181	66/10/90
				cDNA	$\overline{}$	Ι.,		HFIJA29		HFIJA68		HFKES05		HFKEU12		HFPCZ55		HFPDR62		HFPDS07		HFRAB10		HFTBM38		HFTDH56		
				Gene	No.	168		169		170		171		172		173		174		175		176		177		178		

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					Portion		;	31		<del>-</del> -		70		ì					17	,			33		15			
	Last	AA			Pep		- 6	30	,	I3	;	25		,	15				16		<u>8</u>		32		14			
	First Last	AA	Jo	Sig	Pep	<del></del>	,	_		_					-	,	_		_				_		_		-	
	AA	SEQ	А	Ö	<b>&gt;</b>	693	;	694		695		969			269		869		669		90/		701		702		703	
5' NT	of	First	AA of	_	Pep	14		92		163		149			172		258		43		233		280		20		317	
		5' NT	Jo	Start	Codon			92				149			172		258		43		233							
	3, NT	Jo	Clone	Seq.	_	1236		1233		1520		1379			1001		1378		1316		1738		528		1054		1475	
	5' NT 3' NT	Jo	Clone Clone	Seq.		1				9		_			-		-		-		-		-		1		23	
	4,		Total	Ę	Seq.	1236		1233		1520		1379			1001		1378		1316		1738		528		1054		5061	
	Z	SEQ	, П	SO:	X	189		190		191		192			193		194		195		196		197		198		199	
			•		Vector	pBluescript		pBluescript		Lambda ZAP	П	Lambda ZAP	п		Lambda ZAP	П	Lambda ZAP	П	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		pSport1	
		ATCC	Deposit	No:Z	and Date	203960	04/26/99	203960	04/26/99	203960	04/26/99	PTA-	181	66/L0/90	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99
				cDNA	$\overline{}$	HFVGK35		HFVHW43		HFXAV37		HFXBN86			HFXBT66		HFXFZ46		HGBER72		HGBEY14		HGBGN34		HGBHP91		HGCAC19	
			-	Gene	No.	179		180		181		182			183		184		185		186		187		188		189	

		Last	AA	of	ORF	27		38		2		55		24			32		9		327		327		99		5	$\neg$
		First	AA of	Secreted	Portion (	19		59				18		15			11				30		30		36			
	Last	AA	of		Pep	18		28				17		14			10				59		59		35			
	First	AA				1		_		_		-		1			-		_		П		1		1		1	
	AA	SEQ	A	:   	Y	714		715		716		717		718			719		720		721		722		723		724	
5' NT	of	First	AA of	Signal	Pep	156		157		069		62		221			380		453		189		191		1239		61	
		5' NT	ot	Start	Codon			157				62		221					453		189		161		1239		61	
	3, NT		Clone	Seq.		1838		1147		803		1431		1277			531		1093		1976		1982		2142		1009	
	5'  NT 3'  NT	of	Clone Clone	Seq.		1		-		27		1		1			1		1		151		153		1061		1	
			Total	Z	Seq.	1838		1147		1049		1444		1277			531		1093		1980		1982		2154		6001	
	K	SEQ	А	SON.	X	210		211		212		213		214			215		216		217		218		219		220	
					Vector	Uni-ZAP XR		pBluescript		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR			Uni-ZAP XR		Uni-ZAP XR		pBluescript	SK-	pBluescript	SK-	pSport1		pBluescript	SK-
		ATCC	Deposit	No:Z	and Date	096807	04/26/99	096607	04/26/99	203959	04/26/99	203917	04/08/99	PTA-	181	66/L0/90	203960	04/26/99	703960	04/26/99	203960	04/26/99	203960	04/26/99	203959	04/26/99	203957	04/26/99
				cDNA	Clone ID	HHPFU28		HHPSA85		HHSB106		HHSBI65		HHSDI53			HHSFC09		HHSGL28		HILCA24		HILCA24		HISAT67		HJBCU75	
				Gene	No.	200		201		202		203		204			205		506		207		208		509		210	

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		Last	AA	of	ORF	6		27			34		18		43		117		281		50			8		36	
		First	AA of	Sig Secreted	Portion						23				22				21		15			14		19	
	Last	AA	of	Sig	Pep						22				21		10		20		14			13		18	
	First	AA	of	Sig	Pep	-		1			-		-		-		1		-		-			-		-	
	AA	SEQ	П	NO:	Y	725		726			727		728		729		730		731		732			733		734	
5' NT	Jo	First SEQ	AA of	Signal NO:	Pep	527		207			2492		170		256		374		755		207			343		261	
		5' NT	Jo		Codon	•		207											755		207			343		261	
	5' NT 3' NT	Jo	Clone Clone	Seq.	•	599		1017			2886		1298		686		879		1919		1181			1801		2007	
	5' NT	Jo	Clone	Seq.	1	1		1			2233		69		1				581		1			1		1	
			Total	K	Seq.	599		1017			2886		1298		686		628		6161		1181		-	1801		2007	
	N	SEQ	А	NO:	X	221		222			223		224		225		226		227		228			229		230	
					Vector	pCMVSport	3.0	pCMVSport	3.0		pCMVSport	3.0	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		pCMVSport	2.0	pCMVSport	2.0		pCMVSport	2.0	pSport1	
		ATCC	Deposit	No:Z	and Date	203957	04/26/99	PTA-	181	66/10/90	203959	04/26/99	203957	04/26/99	203957	04/26/99	203959	04/26/99	203959	04/26/99	-VIA	181	66/L0/90	203957	04/26/99	203957	04/26/99
				cDNA	Clone ID	HJMAA03		HJMAV41			HJMAY90		HJPBE39		HJPBK28		HJPCH08		HKABU43		HKACI79			HKAFF50		HKGBF25	
				Gene	No.	211		212			213		214		215		216		217		218			219		220	

									5' NT		į	,		!
		ATCC		NI SEO		5' NT 3' NT of of		5' NT	ot First	AA SEO	First AA	Last AA	First	Last
		Deposit		<u>A</u>	Total	Clone Clone			AA of	, A	Jo	Jo	AA of	AA
Gene	cDNA	No:Z		Ö.	N	Seq.	Seq.	Start	Signal	NO:	Sig	Sig	Secreted	of
No.	Clone ID	and Date	Vector	X	Seq.			Codon	Pep	Y	Pep	Pep	Portion	ORF
221	HKIXC44	203957	pBluescript	231	788	343	750	572	572	735	_	76	27	36
222	HKMLK03	203957 04/26/99	pBluescript	232	1049	-	1049	214	214	736	-			Ξ
223	HKMLM95	203957 04/26/99	pBluescript	233	1098	-	1098		390	737	_			4
224	HKTAB41	203957 04/26/99	Uni-ZAP XR	234	797	-	797	172	172	738	-			10
225	HLDBG17	PTA- 181 06/07/99	pCMVSport 3.0	235	652	П	652	184	184	739		23	24	41
226	HLDCA54	203979 04/29/99	pCMVSport 3.0	236	1815	425	1815	550	550	740	_	26	27	46
227	нгрои79	203959 04/26/99	pCMVSport 3.0	237	1488	-	1488	66	66	741	1	23	24	348
227	6/ЛОСТН	203959 04/26/99	pCMVSport 3.0	512	3179	163	1474	75	75	1016	1	29	30	348
228	HLDRT09	203957 04/26/99	pCMVSport 3.0	238	721	254	599	522	522	742	1	20	21	99
229	HLHAP05	203957 04/26/99	Uni-ZAP XR	239	1842	12	1842	45	45	743	1		:	14
230	HLHCS23	203957 04/26/99	Uni-ZAP XR	240	1427	1	1427	25	25	744	1	24	25	34

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		Last	AA	of	ORF	127			7		72		17		$\frac{1}{21}$		10	$\perp$	<del></del>				12			182		_
		First	AA of	Secreted	Portion	47					24	-			70				30		81		12			7		
	Last	AA			Pep	46					23				19				53		17		11					
	First Last	AA	Jo	Sig	Pep	-			_		1		-		_		-									_		
	AA	SEQ	А	NO:	×	745			746		747		748		749		750		751		752		753			754		
5' NT	Jo	First	AA of	Signal	Pep	167			708		441		359		47		214		12		185		305			n		
		5' NT	Jo	Start	Codon	167					441		359		47		214		17		185		305					
	3' NT	of	Clone	Sed.		1768			824		903		926		622		1063		804		268		2450			2385		
	5' NT 3' NT	Jo	Clone Clone	Sed.		1			401		1		1				_				П		-			1652		
			Total	N	Seq.	1768			840		903		926		622		1063		804		268		2450			2385		
	LZ	SEQ	А	NO:	X	241			242		243		244		245		246		247		248		249			250		
					Vector	pCMVSport 1			pCMVSport 1	•	pCMVSport 1		pCMVSport 1		Lambda ZAP	П	Lambda ZAP	П	Lambda ZAP	П	Lambda ZAP	П	Lambda ZAP	п		Lambda ZAP	П	
		ATCC	Deposit	No:Z	and Date	Ι.	792	06/12/60	203957		203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	PTA-	793	09/27/99	PTA-	181	66/L0/90
Ī				cDNA	$\overline{}$	HLIBO72			HLICE88		HLICO10		HLJBS28		HLMBW89		HLMGP50		HLMJB64		HLMMX62		HLQAS12			HLQCL64	,	
				Gene	No.	231			232		233		234		235		236		237		238		239			240		

		Last	AA	 	RH H	52			-	27		57		32		-	 20		43	,	35	-	23		43		14 —	
			AA of	Secreted	Portion	17		19				18		30	-		33		41				12		25			
	Last	AA	of	Sig	Pep	16		18		17		17		53			32		40		27		11		24			
	AA First Last	AA		Sig								Ţ		<b>-</b>			_		1				_		-		<del>-</del>	
	AA	SEQ	_	_	×	755		756		757		758		759			200		761		762		763		764		765	
5, NT	Jo	First	AA of	Signal NO:	Pep	68		192		1751		220		200			122		81		95		363		258		1863	
		5' NT	of	Start	Codon	68		192		1751		220		200			122		81		95				258	1	1863	
	3' NT	_	Clone	Seq.		1243		2564		2488		947		2062			1716		788		1611		626		1146		2966	
	5' NT 3' NT	Jo	Total Clone Clone	Seq.		1		_		1542									-		_		<b>—</b>		Ţ		1527	
			Total	Z	Seq.	1243		2564		2495		947		2062			1716		788		1611		626		1146		2967	
	Ľ	SEQ		SO.	X	251		252		253		254		255			256		257		258		259		260		261	
					Vector	Lambda ZAP	П	pCMVSport	3.0	pCMVSport	3.0	pCMVSport	3.0	pCMVSport	3.0		pCMVSport	3.0	pCMVSport	3.0	pCMVSport	3.0	pSport1		pSport1		pSport1	
		ATCC	Deposit	No:Z	and Date	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	PTA-	795	09/27/60	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203959	04/26/99
				cDNA	$\overline{}$	2		HLWAF06		HLWAU42		HLWAU42		HLWAV47			HLWBB73		HLWCN37		HLWDB73		HLYDF73		HLYEU59		HLYGB19	
				Gene	No.	241		242		243		244		245			246		247		248		249		250		251	

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		Last	AA	_	ORF	73	_	42	450		35		450	_	35	_	35	_	48	_	<u>~</u>	4	<del></del>	_	<u>س</u>	_
		First	AA of	Sig Secreted	Portion	18		21	26		21		56		21		21		18		18					
	Last	AA	of		Pep	17		20	25		20		25		70		70		17		17				19	
	First ]	AA	of	Sig	Pep	-		-	-		-				-		-		_		_		_		_	
	AA	SEQ	А	NO:	X	99/		767	768		69/		2770		771		772		773		774		775		9//	
5' NT	Jo	First	AA of	Signal NO:	Pep	406		211	106		497		106		498		26		211		26		135		30	
		5' NT	of	Start	Codon	406		211	106		497		106		498		26		211		26		135	į	30	
	3, NT	of	Clone	Seq.		752		640	1670		1670		1670		1671		1301		443		1190		1204		2641	
	5' NT 3' NT	Jo	Clone Clone	Seq.		1		-	405		405		406		407		-		_		_		-		1	
			Total	N	Seq.	752		640	1733		1733		1733		1735		1301		443		1190		1204		2641	
	N	SEQ	, П	NO:	X	262		263	264		265		266		267		268		269	:	270		271		272	
					Vector	pSport1		pSport1	Uni-ZAP XR		Uni-ZAP XR		Lambda ZAP	П												
	•	ATCC	Deposit	No:Z	and Date	203957	04/26/99	203957	203917	04/08/99	203917	04/08/99	203917	04/08/99	203917	04/08/99	203917	04/08/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99
		-		cDNA	$\overline{}$	HLYGE16		HLYGY91	HMCAZ04		HMCFH60		HMDAB29		HMDAD44		HMEBB82									
				Gene	No.	252		253	254	· }	255		256		257		258		259		260		261		262	

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		Last			ORF	33		17	1	93	1	31	_	9	$\perp$	32	$\perp$	30			4	27	_	<u>ω</u>	4	381	_
		First	AA of	- 4	Portion	17		13	,	19		23				=		17						56		2	
	Last	AA	of	Sig	Pep	16		12	,	18		22				10		16						25		<u> </u>	
	First	AA	of	Sig	Pep	П		-	,	_		-		-		-		_		-		-		_		<del>-</del>	
	AA	SEQ		NO:	X	777		778		779		780		781		782		783		784		785		186		787	
5' NT	of	First	AA of	Signal NO:	Pep	006		622		113		195		229		1149		249		273		295		120		107	
		5' NT	Jo	Start	Codon	906				113		195		229				249		273		295		120		107	
	3, NT	Jo	Clone	Seq.		2806		2219		1607		1064		1738		1772		2048		799		1396		2945		1667	
	5' NT 3' NT	of	$\circ$	Seq.		884		362		-		_		_		1		-		_		-		-		442	
			Total	NT	Seq.	2836		2276		1607		1064		1738		1772		2048		799		1396		2945		1667	
	N	SEQ	, П	NO:	X	273		274		275		276		277		278		279		280		281		282		283	
					Vector	Lambda ZAP	П	Lambda ZAP	_	ZAP	П	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		pSport1		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR	
		ATCC	Deposit	No:Z	and Date	203957	04/26/99	203957	04/20/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203979	04/29/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203959	04/26/99
				cDNA	$\overline{}$	HMEDE24		HMEDI90		HMELM75		HMIAK10		HMIBF07		HMICI80		HMICP65		HMJAK70		HMSBE04		HMSCL38		HMSCR69	
				Gene	No.	263		264		265		566		267		268		569		270		271		272		273	

					r <sub>T</sub>	<b>—</b>								ı			<u>.                                    </u>										
		1 .	AA	Jo	ORF	66		113	35			48		30		227		9		25		397		38		23	
	<u>.</u>	riisi	AA ot	Secreted	Portion	21		25	12			25		21		26		33		22		2		31		22	
100 1		AA ,		Sig		70		24	11			24		20		25		32		21		-		30		21	
Li trot		<u> </u>	ot	Sig	Pep	1		-	1			-		1		1		1		1		-		1		1	
<	AA	אמע		NO.	Y	788		789	790			791		792		793		794		795		961		797		798	
5' NT	io i	נוואר	AA ot	Signal	Pep	28		20	959			692		710		239		1437		274		137		1015		288	
	7, NT	_		Start	Codon	28	1	50				692		710		239		1437		274		137		1015		288	
3, NT	7 JA 1	, c	Clone	Sed.		1724		2249	2205			3839		2000		2709		2546		1351		2596		2101		1224	
S' NT 3' NT	INI C	5 5		Seq.		1	,	<del></del>	1			-	-	099		1		1327		1		8		927		-	
			•	Ľ	Seq.	1724		2249	2205			3839	ı	2000		2709		2556	•	1351		2596		2288		1224	
LN	CEO	אני אניל		Ö.	X	284		285	286			287		288		289		290		291		292		293		294	
					Vector	Uni-ZAP XR		Uni-ZAP XR	Uni-ZAP XR			pCMVSport	3.0	pCMVSport	3.0	pCMVSport	3.0	pSport1		pSport1	1	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR	
	νΤC	۲) د د	Deposit	No:Z	and Date	203957	04/20/99	203979 04/29/99	PTA-	793	09/27/99	203979	04/29/99	203957	04/26/99	203918	04/08/99	203957	04/26/99	203979	04/29/99	203957	04/26/99	203957	04/26/99	203957	04/26/99
					Clone ID	982HSWH		HMSHU20	HMSHY25			HMTAB77		HMUAE26		HMUAN45		HMVBC31		HMVDU15		HMWBL03		HMWJF53		HNEAK81	
				Gene	No.	274	i	275	276			277		278		279		280		281		282		283		284	

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		Last	AA	of	ORF	34		28		33		32		43		30		32		114		46		59		243		
		First	AA of	Secreted	Portion	24		21		20		23		18		21		24		28		18		23		31		
	First Last	AA	of	Sig		23		20		19		22		17		20		23		27		17		22		30	-	
	First	AA	of	Sig	Pep	1		1		1				1		1		1		1		1		1		1		
	AA	SEQ		ÖN.	Y	66L		008		801		802		803		804		805		908		208		808		608		
5' NT	Jo	First	AA of	Signal	Pep	472		316		02		9/9		314		178		248		89		47		205		237		
		5' NT	Jo	Start	Codon	472		316		02		9/9				178		248								237		
	5' NT 3' NT	of	Clone	Seq.		2710		463		2073		1442		1436		728		915	,	1156		636		1045		1425		
	5' NT	Jo	Total Clone Clone	Seq.	ı	225				1		428		-		1		1		1		1		-		1		
			Total	K	Seq.	2710		489		2073		1442		1436		728		915		1156		989		1045		1425		
	NT	SEQ	А	SO:	X	295		296		297		298		299		300		301		302		303		304		305		
					Vector	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		
		ATCC	Deposit	No:Z	and Date	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	PTA-	181	66/L0/90
				cDNA	Clone ID	HINECL22		HINECW49		HNEDH88		HNFAC50		HINFGR08		HNFHF34		HNGAK51		HNGAM58		HINGBH53		HNGDQ38		HNGDX18		
				Gene	No.	285		286		287		288		589		290		291		292		293		294		295		

ast	AA	of	쥐	132		$\neg$	17		4	$\Box$	22	$\neg$	32		— 89		6			<b>∞</b>	<u> </u>	73			45
First L		Secreted	Portion OKF	19			· <b></b>		25		22		.,		91						+	47			18
			Pep P	18							21	$\dashv$			15						+	46			17
First Last AA AA			Pep						_		1		_		_					-	1	_			
AA SEO	, П	SON.	>	1017			810		811		812		813		814		815			816		817			818
5' NT of First	AA of	Signal NO:	Pep	231			73		58		30		184		181		25			221		252			415
5, NT	Jo	Start	Codon	231							,		184		181							252			415
		Seq.		1411			1002		1103		1029		585		541		1195			1047		1246			1048
5' NT 3' NT of	Clone Clone	Seq.		1			_		_		-		_				_					_			1
	Total	_	Seq.	1411			1002		1103		1029	·	585		541		1195			1047		1246			1048
NT		SON.	X	513			306		307		308		309		310		311			312		313			314
			Vector	Uni-ZAP XR			Uni-ZAP XR			Uni-ZAP XR		Uni-ZAP XR			Uni-ZAP XR										
ATC	Deposit	No:Z	and Date	PTA-	181	66/L0/90	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	PTA-	795	09/27/99	203957	04/26/99	-VLA	181	66/L0/90	203959 04/26/99
		cDNA	Clone ID	HNGDX18		•	HNGDY34		HNGEA34		HINGEQ75		HNGGA68		HNGGP65		69ZHDNH			HNGIV64		HNGJB41			HNGKT41
		Gene	No.	295			296		297		298		299		300		301			302		303			304

		·	NT		5.  NT   3.  NT				AA	_	Last	į	,
	ATCC		SEQ		Jo			First	• .		AA ,	First	Last
	Deposit		A	Total	Total Clone Clone		_	AA of		ot	_	AA of	AA ,
Gene cDNA	No:Z		NO:	Ľ	Seq.	Seq.	Start	Signal	~	Sig		Secreted	ot
Clone ID	and Date	Vector	X	Seq.	_	_	Codon	Pep	Y	Pep	Pep	Portion	ORF
305 HNGMW45	203959	Uni-ZAP XR	315	1530		1530	452	452	819	-	56	27	43
306 HNGNK44		Uni-ZAP XR	316	1178	302	1178	611	611	820	1	18	19	74
307 HNGNO53	203959	Uni-ZAP XR	317	825	-	825	467	467	821	1	15	16	34
308 HNGP125	_	IIni-ZAP XR	318	853	129	853	544	545	822	-	20	21	25
			)   	)									
309 HNHEN82	203918 04/08/99	Uni-ZAP XR	319	573	1	573		28	823	1	13	14	17
310 HNHFE71	203959 04/26/99	Uni-ZAP XR	320	903		903	298	869	824	-			21
311 HNHGK22	203918 04/08/99	Uni-ZAP XR	321	606	-	606	239	239	825	1	26	27	49
312 HNHHB10	203959 04/26/99	Uni-ZAP XR	322	901	1	901	215	215	826	1	28	29	59
313 HNHKS19	203959 04/26/99	Uni-ZAP XR	323	266	1	062	192	192	827	-	26	27	41
314 HNTBT17	PTA- 181 06/07/99	pCMVSport 3.0	324	1959	-	1959	91	16	828				9
315 HNTMH79	203959 04/26/99	pSport1	325	922	1	922	48	48	829	_	35	36	38

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		Last	AA	of	ORF	123		124		123		43		27		33		10		43		71		39		51		
		First	AA of	• 1	Portion	70		70		77		36		∞		18				4		56		25		53		
	Last	AA	of	Sig	Pep	16		19		21		35		7		17				39		25		24		78		
	First	AA	of	Sig	Pep	_		_		_		-	į	-		-		-		-		<del></del>		_		-		
	AA	SEQ		NO:	Y	830		831		832		833		834		835		836		837		838		839		840		
5' NT	of	First	AA of		Pep	148		148		778		43		173		101		248		1714		48		138		230		
		5' NT	of	Start	Codon					778		43				101				1714				138		230		
	3, NT	of	Clone	Seq.		068		892		1249		006		604		1117		927		2216		1356		1036		1349		
	5' NT 3' NT	Jo	Clone Clone	Seq.		1				772		-		1		-		1		1449		$\vdash$		1		-		
			Total	ZZ	Seq.	927		676		1298		006		604		1119		927		2218		1356		9601		1365		
Г	N	SEQ	<sup>'</sup> 白	NO:	X	326		327		328	-	329		330		331		332		333		334		335		336		
					Vector	Uni-ZAP XR		pSport1		pCMVSport	2.0	pCMVSport	2.0															
		ATCC	Deposit	No:Z	and Date	203959	04/26/99	203959	04/26/99	203959	04/26/99	203918	04/08/99	203918	04/08/99	203917	04/08/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	PTA-	795	09/27/99
				cDNA	$\overline{}$	HOABP31		HOABP31		HOACG07		HODAG07		HODBB70		HODBV05		HODCZ32		HOEBK60		HOFAA78		HOFNB74		HOFNU55		
				Gene	No.	316		317		318		319		320		321		322		323		324		325		326		

									5' NT					
				Z		5'  NT 3'  NT	3, NT		of	AA	First	Last		
		ATCC		SEQ		Jo	Jo	5' NT	First	SEQ	AA	AA	First	Last
		Deposit		А	Total	Clone Clone	Clone	Jo	AA of	А	of	of	AA of	AA
Gene	cDNA	No:Z		0 N	N	Seq.	Seq.	Start	Signal	NO:	Sig	Sig	Secreted	of
No.	Clone ID	and Date	Vector	X	Seq.		•	Codon	Pep	Y	Pep	Pep	Portion	ORF
327	HOGBF01	203918	pCMVSport	337	1478	1	1478	309	309	841	1	10	11	20
		04/08/99	2.0						_					
328	HORBS82	203959	Uni-ZAP XR	338	1125	1	1125		21	842	-	19	20	39
		04/26/99												
329	HORBV76	203959	Uni-ZAP XR	339	1157	1	1157	183	183	843	1	25	56	198
		04/26/99												
330	HOSDO75	PTA-	Uni-ZAP XR	340	905	-	305	88	88	844	П			28
		181												
		66/20/90					-							
331	HOSEC25	203959	Uni-ZAP XR	341	1552	1	1552	17	17	845	1	18	19	24
		04/26/99												
332	HOSEI81	203918	Uni-ZAP XR	342	268	1	268	203	203	846	-	22	23	83
		04/08/99												
333	HOSEJ94	203979	Uni-ZAP XR	343	1767	622	1750	848	848	847	1	21	22	28
		04/29/99												
334	HOUCA21	203918	Uni-ZAP XR	344	1129	1	1129	200	200	848	_	27	28	33
		04/08/69											•	
335	HOUDE92	203918	Uni-ZAP XR	345	1284	-	1282		70	849	-	9	7	88
		04/08/99												
336	HOUDR07	203959	Uni-ZAP XR	346	1161	1	1911	170	170	850	1	27	28	65
		04/26/99												

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Last	AA	of	ORF	11			72		20			26		25		30		66		45		19			17		
First	AA of	Secreted	Portion				43					20				24		14		40		18					
First Last AA AA	Jo	Sig	Pep				42					19				23		13		39		17					
		Sig	Pep	1			1		1			-		-		1		1		1		1			1		
AA SEQ	D	Ö.	Y	851			852		853			854		855		856		857		828		859			988		
5' NT of First	AA of	Signal NO:	Pep	144			520		188			252		184		1021		188		258		94			38		
5' NT	Jo		Codon				520		188			252		184		1001		188		258		94			38		
5' NT 3' NT of of		Seq.		662			2882		1102			1129		2587		2002		585		835	·	879			352		
5' NT of	Clone	Seq.		92			457		45			1	,	1		803		1		Ţ		1			1		
			Seq.	833			2927		1249		- "	1129		2587		3097		585		835		628 -			352		
NT SEQ		NO:	X	347			348		349			350		351		352		353		354		355			356		
			Vector	Uni-ZAP XR			Uni-ZAP XR		Uni-ZAP XR			pSport1		Uni-ZAP XR			Uni-ZAP XR										
ATCC	Deposit	No:Z	and Date	PTA-	181	66/1/0/90	203959	04/26/99	PTA-	793	09/27/99	203918	04/08/66	203918	04/08/69	203917	04/08/99	203959	04/26/99	203959	04/26/99	PTA-	181	66/1/0/90	PTA-	181	66/L0/90
		cDNA	Clone ID	HOUED72			HOUFS04		HOUHI25			HOVBD85		HPCAB41		HPCAL26		HPEAD23		HPFBA54		HPFCI36			HPFDI37		
		Gene	No.	337			338		339			340		341		342		343		344		345			346		

									5' NT					
				Ż		5' NT 3' NT	3, NT		Jo	AA	First Last	Last		
		ATCC		SEQ		of	Jo	5' NT	First	SEQ	AA	AA	First	Last
		Deposit			Total	Clone Clone	Clone	ot	AA of	А			AA of	AA
Gene	cDNA	No:Z		NO:	Ľ	Seq.	Seq.	Start	Signal	:     	Sig	Sig	Secreted	of
No.	Clone ID	and Date	Vector	X	Seq.	'	_	Codon	Pep	Y		Pep	Portion	ORF
347	HPIAA80	203959 04/26/99	Uni-ZAP XR	357	616	312	616		314	861	-	13	14	37
348	HPJBJ51	203959 04/26/99	Uni-ZAP XR	358	2793	522	2421	715	715	862	1	14	15	69
349	HPJBJ51	203959 04/26/99	Uni-ZAP XR	329	2795	523	2422	716	716	863	1	14	15	69
350	HPJBU43	PTA- 181	Uni-ZAP XR	360	575	-	575		242	864	1			17
		06/0/90												
351	HPJCW58	203918	Uni-ZAP XR	361	1165	-	1165	177	177	865	-	19	70	28
352	HPMBX22	203959	Uni-ZAP XR	362	454	-	454		211	998	1			19
353	HPMCJ84	203918	Uni-ZAP XR	363	788	1	788	83	83	867	-	22	23	38
354	HPMCV30	203918 04/08/99	Uni-ZAP XR	364	806	-	806	52	52	868	1	27	28	47
355	HPMFH77	203918 04/08/99	Uni-ZAP XR	365	1891	-	1891		251	869	1	11	12	35
356	HPQAX38	203979 04/29/99	Lambda ZAP II	396	1157	41	1157		295	870	-	10	11	16
357	HPQAX38	203979 04/29/99	Lambda ZAP II	367	1158	41	1158		295	871	1	10	11	16

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		Last	AA	of	ORF	34		35		134		39		35			08		13		76		25			4	
		First		9,	Portion	31		34		19		23		17			17						70			25	
	Last	AA	Jo	Sig	Pep	30		33		18		22		16			16						19			24	
	First ]	AA	of	Sig	Pep	_		_				_		-			_		-		1		<u> </u>			_	
	AA	SEQ	А	0 N	χ	872		873		874		875		928			877		878		879		880			881	
2, NL	Jo	First SEQ	AA of	Start Signal	Pep	85		16		684		1810		265			885		194		405		220			122	
		5' NT	of	Start	Codon	85		16		684		1810				i	885		194		405						
	3, NT	of	Clone	Seq.		2267		434		1648		2757		689			1680		325		854		1496			1135	
	5' NT 3' NT	Jo	Total Clone Clone	Seq.		1		-		558		1701		_			658		_		240		1			1	
			Total	N	Seq.	2267		434		1673		2805		709			1760		325		878		1496			1135	
	Z	SEQ		NO:	X	368		369		370		371		372			373		374		375		376			377	
					Vector	Lambda ZAP	П	Lambda ZAP	П	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR			pBluescript		Uni-ZAP XR		Uni-ZAP XR		pCMVSport	3.0		pCMVSport	3.0
		ATCC	Deposit	No:Z	and Date	203918	04/08/99	203918	04/08/99	203959	04/26/99	203959	04/26/99	PTA-	181	66/10/90	203959	04/26/99	203918	04/08/99	203959	04/26/99	PTA-	181	66/L0/90	203959	04/26/99
				cDNA	$\overline{}$	HPQCB83		HPQCC53		HPRBH85		HPRCA64		HPRCD35			HPTRM02		HPWBA29		HPWDK06		HRAAD30			HRADA42	
				Gene	No.	358		359		360		361		362			363		364		365		366			367	

		+			山	~		$\neg$		Т	-	П		$\neg$						$\neg$	_	Т	_				$\neg$
		Last	AA	Jo	ORF	253			65		63		31		46		6		72		10		10		33		
		First	AA of	Sig Secreted	Portion	40			18		53		28		18				21						56		
	Last	AA	of	Sig	Pep	39			17		78		27		17	ļ			20						25		
	AA First Last	AA	of	Sig	Pep	-			_		1		-		-		П				_				-		
	AA	SEQ	А	ÖN N	Y	882			883		884		885		988		887		888		889		890		891		
5' NT	Jo	First	AA of	Signal NO:	Pep	691			198		233		578		215		32		649		120		1958		50		
		5° NT	of	Start	Codon	169			198		233		578						649		120				20		
	3, NT		Clone	Seq.		2684			1206		1324		1500		773		543		1638		726		2271		281		
	5' NT 3' NT	of	Total Clone Clone	Seq.	•	1			17		_		547				1		711				1687		1		
	•			Z	Seq.	2704			1225		1324		1500		<i>9LL</i>		543		1681		728		2301		281		
	Z	SEQ	А	Ö.	X	378			379		380		381		382		383		384		385		386		387		
		•			Vector	pCMVSport	3.0		pCMVSport	3.0	pCMVSport.	3.0	Uni-ZAP XR		Uni-ZAP XR	,	Uni-ZAP XR										
		ATCC	Deposit	No:Z	and Date	PTA-	181	66/L0/90	203959	04/26/99	203959	04/26/99	203918	04/08/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	PTA-	181	66/10/90
				cDNA	$\overline{}$	HRADF49			HRADN25		HRADT25		HRDAI17		нкрродз9		HRDER22		HRDEX93		HRDFK37		HRGBD54		HROEA08		
				Gene	No.	368			369		370		371		372		373		374		375		376		377		

		Last	AA	Jo	ORF	56		22		63		37		98		98		10		13		16		17		0/	
		First	AA of	Secreted	Portion	18		17		30				17		16		8				20		12		21	
	Last	AA	of	Sig	Pep	17		16		59				16		15		7				19		11		20	
	First	AA	of	Sig	Pep	1		1		1		1		1		1		1		1		1		1		1	
	AA	SEQ			Y	892		893		894		895	,	968		268		868		668		006		901		905	
2, NT	Jo	First	AA of	Signal NO:	Pep	99		129		159		124		106		445		129		304		257		473		249	
		5' NT	of	Start	Codon			129				124		106		445				304		257				647	
	3, NT	Jo	Clone	Seq.	ı	1061		265		349		6101		214		554		1213		882		1648		762		1474	
	5' NT 3' NT	Jo	Clone Clone	Seq.	•	1		1		1		1		1		1		1		I		1		1		452	
			Total	NT	Seq.	1001		565		349		1019		214		554		1273		882		1648		<i>1</i> 62		1474	
	NT	SEQ		NO:	X	388		389		390		391		392		393		394		368		396		397		868	
					Vector	Uni-ZAP XR		pBluescript		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR									
		ATCC	Deposit	No:Z	and Date	203918	04/08/69	503959	04/26/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	503959	04/26/99	203959	04/26/99	503959	04/26/99	203918	04/08/99	203918	04/08/99	203979	04/29/99
				cDNA	Clone ID	HSAVA08		HSAVW42		HSAWN53		HSAWZ40		HSAYC41		HSDZM54		HSHBF76		HSIFG47		HSJBY32		HSKDR27		HSLHG78	
				Gene	No.	378		379		380		381		382		383		384		385		386		387		388	

_					_													_								_	
		Last	AA	Jo	ORF	41		4		14		62		42		99		11			265		21			21	
		First	AA of	Secreted	Portion	21						15		12		29					15						
	Last	AA	Jo	Sig	Pep	20						14		11		28					14						
	AA First Last	AA	Jo	Sig	Pep	1		1		-		1		-		1					-		1			1	
	AA	SEQ	А	NO:	Y	903		904		905		906		706		806		606			910		911			912	
5' NT	Jo	First	AA of	Start Signal	Pep	485		941		164		1508		206		229		133		·	195		128			253	
		5' NT	of	Start	Codon	485												133			195					253	
	3, NT	of	Total Clone Clone	Seq.		655		1286		979		1765		721		1024		1210			1428		1633			1406	
	5' NT 3' NT	of	Clone	Seq.		1		735		1		1391		_		1		1		•	1012		13				
			Total	N	Seq.	655		1286		979		2186		721		1024		1210			1445		1633			1406	
	N	SEQ	Ω	NO:	X	399		400		401		402		403		404		405			406		407			408	
					Vector	Uni-ZAP XR	:	Uni-ZAP XR		Uni-ZAP XR			Uni-ZAP XR		Uni-ZAP XR			Uni-ZAP XR									
		ATCC	Deposit	No:Z	and Date	503959	04/26/99	203959	04/26/99	203918	04/08/69	203959	04/26/99	203959	04/26/99	203918	04/08/99	-VLA	181	66/20/90	203959	04/26/99	PTA-	791	09/27/99	203918	04/08/99
				cDNA	Clone ID	HSLHX15		HSNAP85		HSNAZ09		HSNBM34		HSOAH16		HSQBF66		HSQDO85			HSQES57		HSRBE06			HSSDI26	
				Gene	No.	389		330		391		392		393		394		395			396		397			398	

				ſŢ.											_										_		$\neg$
	Last	AA		ORF	62			09		125		38		59		14			33		27		4		137		
	First	AA of	Secreted	Portion	<i>L</i> 1			56		20		18		32					22		20				11		
I act	AA	of		Pep	16			25		19		17		34					21		19				91		
Firet I act			Sig	Pep	1			1		1		1		1		1			_		1		1		1		
AA	SEQ	П	NO:	Y	913			914		915		916		917		918			919		920		921		925		
5° NT of	14	AA of	Signal NO:	Pep	28			184		264		245		380		211			232		396		306		39		
	5' NT	Jo	Start	Codon	28							245	;	380					232		366						
3, NT		Total Clone Clone	Seq.		1274			1053		1133		1954		710	·	2206			926		1198		2174		1764		
ty's'Nt	of Jo	Clone	Seq.		1			1	·	85		1		250		1			_		_		1		1		
		Total	Z	Seq.	1282			1053		1238		1954	·	874		2206			926		1198		2174		1764		
ĘŻ	SEQ	O	NO:	X	409			410		411		412		413		414			415		416		417		418		
				Vector	Uni-ZAP XR			Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		pCMVSport	3.0	Uni-ZAP XR			pCMVSport	3.0	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		
	ATCC	Deposit	No:Z	and Date	PTA-	181	66/L0/90	203959	04/26/99	203959	04/26/99	203918	04/08/99	503959	04/26/99	PTA-	795	09/27/99	203918	04/08/99	503959	04/26/99		04/26/99	PTA-	181	66/L0/90
			cDNA	Clone ID	HSSEA64			HSSEF77		HSSFE38		HSSGJ58		HSWBE76		HSXCP38			HSYBI06		HT1SC27		HT3BF49		HT4FV41		
			Gene	No.	399			400		401		402		403		404			405		406		407		408		

	1	Last	AA	of	ORF	50		31		4		20		22		99		23			5		45		36		32	
	-					5		~		_		2		2		2	-	2					4		3		3	$\dashv$
	į	First	AA of	Secreted	Portion	18		24						21		23		20					61		25		20	
_		AA	of	Sig	Pep	<b>L</b> I		23						20		77		61					18		24		61	
į	First	-	Jo	Sig	Pep	1		1		I		1		1		1		Ţ			1		I		I		1	
•	AA	- 4	А	NO:	Y	923		924		925		976		927		928		676			930		931		932		933	
5' NT	t of	First	AA of	Signal NO:	Pep	228		135		632		151		1017		961		211			325		287		760		261	
		$\sim$	Jo	Start	Codon			135		632		151		1017		961		211			325				760		197	
			Clone	Seq.		682		1743		1623		825		2221		1662		2055			829		1247		1587		2179	
E.	5. NI 3. NI	ō	Clone Clone	Seq.		59						1		57		106		-			1		1		1		1	
			Total	N	Seq.	682		1743		1623		825		2221		1662		2055			829		1247		1587		2179	
Ę		SEQ DES	А	NO:	X	419		420		421		422		423		424		425			426		427		428		429	
					Vector	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		pSport1		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR			Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR	
	Č	ATCC	Deposit	No:Z	and Date	203959	04/26/99	203959	04/26/99	203918	04/08/99	203918	04/08/99	203959	04/26/99	203959	04/26/99	PTA-	181	66/20/90	203959	04/26/99	203959	04/26/99	203918	04/08/99	203959	04/26/99
				cDNA	Clone ID	HT5FX79		HT5GR59		HTAEI78		HTDAA78		HTEAG62		HTECB02		HTECC15			HTEDF18		HTEDJ28		HTEDS12		HTEED26	
				Gene	No.	409		410		411		412		413		414		415			416		417		418		419	

													-									ı					
		Last	AA	of	ORF	32		7		_		323		37		7			88		34		23		38		
		First	AA of	Secreted	Portion	20						31		21					9		20				25		
	Last	AA	of	Sig	Pep	19						30		20					S		19				24		
	First Last	AA	of	Sig	Pep	_		1		_		_		-							1		1		_		
	AA		О	Ö	Y	934		935		936		937		938		939			940		941		942		943		
S' NT	Jo	First SEQ	AA of	Start   Signal NO:	Pep	259		262		262		182		493		173			280		170		101		171		
		5° NT	of	Start	Codon	259		262		262		182									170		101		171		
	5' NT 3' NT		Total Clone Clone	Seq.		2159		984		984		1263		908		981			1400		1504		1324		2116		
	5' NT	Jo .	Clone	Sed.		1		45		45		110				1			529		1		1		1		
			Total	N	Seq.	2167		1015		1273		1282		908		981			1402		1523		1324		2116		
	Z	SEQ	П	SON:	X	430		431		432		433		434		435			436		437		438		439		
					Vector	Uni-ZAP XR		Uni-ZAP XR			Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		!								
		ATCC	Deposit	No:Z	and Date	203959	04/26/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	PTA-	181	66/L0/90	203959	04/26/99	203959	04/26/99	203959	04/26/99	PTA-	181	66/20/90
				cDNA	$\overline{}$	HTEED26		HTEEF26		HTEEF26		HTEEW69		HTEGS07		HTEGS11			HTEHA56		HTEHU59		HTEJD29		HTEKM46		
				Gene	No.	420		421		422		423		424		425			426		427		428		429		

				Ę			1		5° NT	× ×	1	1 004		
				Z		5. NT 3. NT			IO	AA	FIFS	Last	-	•
		ATCC		SEQ		of		5' NT	First	• .	AA	AA	First	Last
		Deposit		А	Total	Clone Clone	Clone	of	AA of	А	of		AA of	AA
Gene	cDNA	No:Z		0 N	Z	Seq.	Seq.	Start	Signal	~		-		Jo
So.	Clone ID	and Date	Vector	X	Seq.			Codon	Pep	≻	Pep	Pep	Portion	ORF
430	HTEMQ17	203959	Uni-ZAP XR	440	1768	1	1768	446	446	944	-			12
		04/26/99					,		3	1	1	1	5	1
431	HTENR63	PTA-	Uni-ZAP XR	441	1591		1591	132	132	945		22	77	ဂ္ဂ
		792												
		09/27/99										1	,	
432	HTGGM44	203959	Uni-ZAP XR	442	3016		2761	179	179	946				<b>8</b>
		04/26/99												
433	HTHBZ06	203959	Uni-ZAP XR	443	623	193	619	318	318	947	_			
		04/26/99												
434	HTLAP64	203918	Uni-ZAP XR	444	1092	_	1092	173	173	948		19	70	70
		04/08/99										$\neg$		1
435	HTLBT80	203959	Uni-ZAP XR	445	2101	817	1881	912	912	949	_	27		129
		04/26/99												1
436	HTLDA84	203918	Uni-ZAP XR	446	1444	_	1444		225	950	<del>-</del>			13
		04/08/99												
437	HTLDN29	203959	Uni-ZAP XR	447	1374	-	1348	175	175	951	_	23	24	33
		04/26/99												
438	HTLDU78	203918	Uni-ZAP XR	448	1318	-	1318	219	219	952	_			∞
		04/08/99	:											
439	HTLEC82	203959	Uni-ZAP XR	449	1260	217	11119	530	530	953	_	34	35	36
		04/26/99												,
440	HTLEM16	203959	Uni-ZAP XR	450	1915	1158	1755	1220	1220	954		27	28	69
		04/70/99												

		Last	AA	Jo	ORF	207		6		31		23		37		38		75		75		75		75		75	
		First L	AA of	Secreted	Portion C	31   2			1	6			1	17		34		70	1	 20		70		70		70	
	Last	AA	of	Sig	Pep	30			1	∞				16		33		19		19		19		19		19	
	First Last	AA	of	Sig	Pep			-		-		_		-		_		_		1		_				-	
	AA	SEQ	А		Y	955		1018		926		957		928		959		096		961		962		696		964	
5, NT	Jo	First	AA of	Signal NO:	Pep	205		91		209		340		1802		933		642		644		644		644		644	
		5' NT	of	Start	Codon	205		91				340		1802		933		642		644		644		644		644	
	3, NT		Clone	Seq.		1070		1065		1160		1159		2377		1968		1100		1033		1033		1033		1033	
	5' NT 3' NT	Jo	Clone Clone	Seq.		1		1				_		1205		098		140		142		142		142		142	
			Total	K	Seq.	1070		1065		1160		1159		2377		1968		1100		1081		1044		1081		1081	
	Z	SEQ	A	SO.	X	451		514		452		453		454		455		456		457		458		459		460	
					Vector	Uni-ZAP XR																					
		ATCC	Deposit	No:Z	and Date	203918	04/08/99	203918	04/08/66	203918	04/08/99	203979	04/29/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	203959	04/26/99
				cDNA	$\overline{}$	HTLEV48		HTLEV48		HTLFA13		HTLFI73		HTLGI89		HTLIF11		HTLIF12									
				Gene	No.	441		441		442		443		444		445		446		447		448		449		450	

									5' NT					
				N		5'  NT 3'  NT	3, NT		of	AA	First Last	Last		
		ATCC		SEQ		of	Jo	5' NT	First	SEQ	AA	AA	First	Last
		Deposit			Total	Total Clone Clone	Clone	of	AA of	А	Jo	oę	AA of	AA
Gene	cDNA	No:Z		NO:	M	Seq.	Seq.		Signal	NO:	Sig	Sig	Secreted	Jo
No.	_	and Date	Vector	X	Seq.	•		Codon	Pep	Y	Pep	Pep	Portion	ORF
451	HTLIF12	203959	Uni-ZAP XR	461	1801	142	1033	644	644	965	П	19	70	75
		04/26/99												
452	HTNAM63	203918	pBluescript	462	1006	1	1006		193	996	_	15	16	30
		04/08/66	SK-											
453	HTNBK13	203959	pBluescript	463	1160	295	1148	534	534	196	_	16	17	21
		04/26/99	SK-											
454	HTOAI50	203959	Uni-ZAP XR	464	1258		1258	19	61	896		17	18	27
		04/26/99												
455	HTOAM11	203918	Uni-ZAP XR	465	1200	1	1200	68	68	696	_	24	25	34
		04/08/99												
456	HTODH57	203918	Uni-ZAP XR	466	1652	1	1652		228	026	-	18	19	71
		04/08/99							,					
457	нторн83	203918	Uni-ZAP XR	467	1981		1981	103	103	971	1	21	22	32
		04/08/99												
458	HTOEV16	PTA-	Uni-ZAP XR	468	1640		1640	201	201	972	П	39	40	118
		181												
		66/L0/90												
459	HTOGR38	203959	Uni-ZAP XR	469	9//	138	9/1		314	973	_	23	24	42
		04/26/99					-							
460	HTOH021	203918	Uni-ZAP XR	470	727	I	727		439	974		2	9	63
	-	04/08/99												

			Ĺ		S' NIT 3' NIT	Z, NT		5' NT of	<b>V V</b>	Firet I get	I act		
	ATCC		SEQ		of		5' NT		SEQ	AA	AA	First	Last
	Deposit		А	Total	Clone Clone	Clone	Jo	AA of	А		of	AA of	AA
Gene cDNA	No:Z		NO:	N	Seq.	Seq.	Start	Signal	Ö	Sig	Sig	Secreted	ot
)	and Date	Vector	X	Seq.			Codon	Pep	Y	Pep	Pep	Portion	ORF
471 HTXDB22	PTA-	Uni-ZAP XR	481	1211	1	1135		229	586	1	10	=	22
	181							•					
	66/L0/90									Í			
472 HTXDC38	203959	Uni-ZAP XR	482	820	106	908	359	359	986	_		•	18
	04/26/99												
473 HTXDC77	203979	Uni-ZAP XR	483	1441	159	1400	99	65	286	1	18	19	151
	04/59/99												
474 HTXDD61	PTA-	Uni-ZAP XR	484	1140	1	1140		49	886	-	17	18	132
	181												
	66/L0/90												
475 HTXDG92	203959	Uni-ZAP XR	485	1162	1	1162		216	686		24	25	99
	04/26/99												
476 HTXET11	203918	Uni-ZAP XR	486	686	_	686	178	178	066	Н	22	23	53
	04/08/99												
477 HTXFA72	PTA-	Uni-ZAP XR	487	1861	1	1861	192	192	166		17	18	29
	181												
	66/L0/90												
478 HTXJY08	203959	Uni-ZAP XR	488	1187	12	1187	108	108	665	1			16
	04/26/99												
479 HTXKF95	203959	Uni-ZAP XR	489	884	6/	875	330	330	666	1	28	29	78
	04/26/99												
480 HTXMZ07	203959	Uni-ZAP XR	490	1652	189	1640	319	319	994	_	22	23	37
	04/26/99												

									5' NT					
				N		5' NT 3' NT	3, NT		Jo	AA	First Last	Last		
		ATCC		SEQ		of		5' NT	First	SEQ	AA	AA	First	Last
		Deposit		Α	Total	$\circ$	Clone	of	AA of	А		of	AA of	AA
Gene	cDNA	No:Z		NO:	IN	Seq.	Seq.	Start	Signal NO:	: ON	Sig		• .	of
No.	Clone ID	and Date	Vector	X	Seq.			Codon	Pep	X	Pep	Pep	Portion	ORF
481	HUFCL31	203959	pSport1	491	1460	1	1460		287	995	_			56
		04/26/99												
482	HUKBT67	203959	Lambda ZAP	492	2069	74	2052		273	966	<b>—</b>	21	22	39
		04/26/99	П											
483	HUKDF20	203918	Lambda ZAP	493	1105		1105	214	214	266	_	70	21	33
		04/08/99	П											
484	HUKDY82	203918	Lambda ZAP	494	1435	-	1435	187	187	866		17	18	32
		04/08/99	П											
485	HUSCJ14	PTA-	Lambda ZAP	495	3342		3342	74	74	666	_	30	31	196
		1838	П											
		02/06/00												
486	HUSGL67	203918	pSport1	496	1008	65	1008	350	350	1000		21	22	47
		04/08/68												
487	HUSGU40	203959	pSport1	497	1054	_	1054		200	1001		70	21	46
		04/26/99												
488	HUSIR18	203959	pSport1	498	9/8	<b>—</b>	928	83	83	1002	_	16	17	22
		04/26/99												
489	HUVDJ48	203918	Uni-ZAP XR	499	1827		1827	196	196	1003				S
		04/08/99												
490	HWAAI12	203959	pCMVSport	200	3303	1	1838	223	223	1004	_			53
		04/26/99	3.0											
491	HWBBQ70	203959	pCMVSport	501	1948	1	1948	222	222	1005	-	21	22	43
		104/70/33	3.0											

									5' NT					
			•	N		5' NT 3' NT	3, NT		of	AA	AA First Last	Last		
	•	ATCC		SEQ		of	of	5' NT	First SEQ	SEQ	AA	AA	First	Last
		Deposit			Total	Clone	Total Clone Clone	of	AA of	А	of	ot	AA of	AA
Gene	cDNA	No:Z		NO:	Z	Seq.	Seq.	Start	Start Signal NO:		Sig	Sig		Jo
No.	Clone ID	and Date	Vector	×	Seq.		-	Codon	Pep	Y	Pep	Pep	Portion	ORF
492	HWBCN36	203959	pCMVSport	505	1008	1	1008	378	378	1006		23	24	96
		04/26/99	3.0											
493	HWBDJ08	203959	pCMVSport	503	2085		2085	253	253	1007	_	59	30	20
		04/26/99	3.0											
494	HWBFX16	203959	pCMVSport	504	1497	1	1497		267	1008	_			3
		04/26/99	3.0											
495	HWDAC26	203959	pCMVSport	205	1958	1	1958	242	242	1009		25	56	35
		04/26/99	3.0						3					
496	HWDAG96	203959	pCMVSport	909	1147	300	1147	998	998	1010	_	18	19	32
		04/26/99	3.0											
497	HWDAJ01	203959	pCMVSport	202	781	—	781	288	288	1011	_			75
		04/26/99	3.0											
498	HWHPB78	203959	pCMVSport	508	1346	_	1346	700	200	1012	1	23	24	99
		04/26/99	3.0											
499	HYABC84	203959	pCMVSport	209	1338	292	1238	1015	1015	1013	<del></del>	28	53	62
		04/26/99	3.0											
200	HYABC84	203959	pCMVSport	510	1478	833	1306	1080	1080   1014	1014		28	- 50	62
		04/26/99	3.0											

# D9950D82.091201

#### TABLE 1B

		_			AA		Tissue Distribution		OMIM
Clone ID   Contig   SEQ ID   ORF   SEQ	SEQ ID ORF	ORF	,	SEQ			Library code: count	Cytologic	Disease
ID: NO: X (From-To)	NO: X (From-To)	(From-To)		UO: Y		Predicted Epitopes	(see Table 4 for Library Codes)	Band	Reference(s):
H6BSF56 762968 11 83 - 508 515	11 83 - 508	83 - 508	- 508	515	I	Asn-131 to Met-140.	AR089: 53, AR060: 29 L0599: 4, L0439: 3, L0777:		
							3, H0253: 2, H0520: 2,		
							L0754: 2, L0745: 2, L0759:		
							2, H0556: 1, H0657: 1, S0116: 1 H0450: 1 S0418:		
							1 50046: 1 50220: 1, 30416:		
							1, 50040: 1, 50222: 1, H0492: 1, S0049: 1, H0123:		
							1, H0050: 1, H0051: 1,		
							H0615: 1, S0036: 1, H0494:		
	-	_					1, L0805: 1, L0776: 1,		•
							S0126: 1, H0435: 1, H0670:		
							1, S0028: 1, L0747: 1,		
							S0026: 1 and H0542: 1.		
H6EDM64 841331 12 1448 - 1468 516	12 1448 - 1468	1448 - 1468	- 1468	516	⊢-		AR060: 22, AR089: 16		
					_		H0333: 6, H0556: 5,		
							H0255: 5, H0547: 5, H0618:		
							4, H0581: 4, H0553: 4,		
							H0135: 4, L0783: 4, S0358:		
-							3, S0222: 3, H0318: 3,		
							H0052: 3, H0617: 3, L0769:		
							3, H0521: 3, H0555: 3,		
							H0436: 3, H0423: 3, H0341:		
							2, H0402: 2, H0619: 2,		
		-					H0549: 2, H0592: 2, H0253:		
							2, S0474: 2, H0620: 2,		
							H0181: 2, H0059: 2, H0561:		
							2, L0761: 2, L0764: 2,		
							L0809: 2, H0520: 2, H0682:		
	-	-					2, S0330: 2, H0522: 2,		

7: 2, L0750: 596: 2, 4: 1, H0686: 6049: 1, 66: 1, H0484: 356: 1, 4: 1, S0360: 300: 1, 6: 1, H0370: 5586: 1, 59: 1, L0623: 6082: 1, 59: 1, H0424: 6082: 1, 60: 1, H0633: 6487: 1, 60: 1, H0633: 648: 1, 60: 1, L0774: 61: 1, L0796: 648: 1, 62: 1, L0774: 63: 1, S0126: 770: 1, 77: 1, S0028: 780: 1, 77: 1, S0011: 196: 1 and	47: 4, L0794: 046: 2, 9: 2, H0575:
L0751: 2, L0747: 2, L0750: 2, L0755: 2, L0596: 2, L0601: 2, H0624: 1, H0686: 1, H0295: 1, T0649: 1, H0657: 1, H0656: 1, H0686: 1, H0657: 1, H0656: 1, H0370: 1, S0442: 1, S0356: 1, S0342: 1, S0342: 1, S0342: 1, S0342: 1, S0342: 1, S0342: 1, H0457: 1, H0486: 1, H0589: 1, H0587: 1, H0486: 1, H0687: 1, H0618: 1, H0606: 1, H0424: 1, H0618: 1, H0606: 1, H0424: 1, H0618: 1, H0606: 1, H0494: 1, H0664: 1, H0494: 1, H0624: 1, H0494: 1, H0625: 1, H0130: 1, H0633: 1, H0647: 1, S0426: 1, H0647: 1, S0426: 1, H0647: 1, S0426: 1, L0664: 1, L0669: 1, L0678: 1, H0647: 1, S0027: 1, H0647: 1, S0028: 1, H0647: 1, S0027: 1, S0028: 1, L0678: 1, H0448: 1, S0011: 1, H0136: 1, S0196: 1 and 1, H0136: 1, S0196: 1 and	H0352: 1. L0809: 4, L0747: 4, L0794: 3, L0759: 3, S0046: 2, H0497: 2, H0559: 2, H0575:
	517
	263 - 319
	13
	889401
	H6EEC72
	3

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2, H0618: 2, H0050: 2, L0769: 2, L0766: 2, L0663: 2, H0521: 2, L0743: 2, L0748: 2, H0520: 1, H0550: 1, H0650: 1, H0650: 1, H0657: 1, H0655: 1, H0650: 1, H0650: 1, H0635: 1, H0635: 1, H0635: 1, H0635: 1, H0640: 1, H0635: 1, H0069: 1, H0646: 1, H0647: 1, L0770: 1, L0770: 1, L0644: 1, L0775: 1, L0666: 1, L0665: 1, H0651: 1, L0665: 1, H0659: 1, H0659: 1, L0755: 1, L0665: 1, L0755: 1, L0665: 1, L0755: 1, L0755: 1, L0755: 1, L0755: 1, L0755: 1, H0422: 1, H0420: 1, H0422: 1, H0422: 1, H0422: 1, H0420: 1, H04	L0748: 4, H0457: 3 and S6022: 1.	AR251: 7, AR310: 6, AR265: 6, AR053: 6, AR060: 6, AR055: 5, AR312: 5, AR309: 5, AR273: 5, AR061: 5, AR206: 5, AR194: 5, AR186: 5, AR213: 4, AR052: 4, AR089: 4, AR253: 4, AR248: 4, AR253: 4, AR248: 4, AR205: 4, AR033: 3, AR243: 3, AR096: 3, AR104: 3, AR26: 3,
	Leu-6 to Ser-12.	Arg-14 to Ile-24.
	518	519
	135 - 371	250 - 327
	14	15
	584773	847112
	HACAB68	HACBJ56
	4	٥

AR263: 3, AR204: 2, AR244: 1, AR249: 1 H0661: 1, S0045: 1, H0550: 1, S0280: 1, S0010: 1, H0028: 1, L0764: 1, L0803: 1, L0665: 1, S0053: 1, H0670: 1, L0748: 1, L0731: 1 and L0581: 1.	H0052: 6, S0002: 5, H0580: 3, S0051: 3, L0766: 3, L0439: 3, L0777: 3, L0361: 3, S0046: 2, H0619: 2, H0550: 2, S0280: 2, H0039: 2, S0142: 2, L0794: 2, L0775: 2, L0748: 2, L0754: 2, L0747: 2, L0758: 2, L0756: 2, H0170: 1, H0265: 1, H0556: 1, S0040: 1, H0661: 1, H0663: 1, S0420: 1, S0356: 1, S0354: 1, H0641: 1, H0540: 1, H0431: 1, H0586: 1, H0492: 1, H0431: 1, H0548: 1, H0545: 1, H0053: 1, S0476: 1, H0545: 1, H0053: 1, H0042: 1, H0053: 1, H0042: 1, L0703: 1, H0561: 1, S0372: 1, S0456: 1, L0770: 1, L0766: 1, L0770: 1, L0769: 1, L0662: 1, H0539: 1, H0521: 1, L0770: 1, L0762: 1, L0770: 1, L0772: 1, L0770:
	Cys-2 to Leu-8.
	220
	217 - 342
	16
	847113
	HACBS22
	φ

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1, H0668: 1 and H0506: 1.	AR089: 25, AR060: 15 L0759: 6, L0769: 5, H0052: 4, L0770: 4, L0809: 4, L0439: 4, L0752: 4, S0408: 3, L0751: 3, L0747: 3, L0779: 3, S0007: 2, H0351: 2, H0333: 2, H0427: 2, H0581: 2, L0662: 2, L0649: 2, L0774: 2, L0806: 2, L0777: 2, H0541: 2, L0777: 2, H0543: 1, H0723: 1, H0171: 1, H0575: 1, H0722: 1, H0733: 1, S0140: 1, H0125: 1, H0592: 1, H0549: 1, H0587: 1, H0587: 1, H0586: 1, H0678: 1, H056: 1, H0123: 1, T0010: 1, H0266: 1, H0673: 1, S0364: 1, H0673: 1, L0772: 1, L0646: 1, L0761: 1, L0772: 1, L0646: 1, L0761: 1, L0772: 1, L0646: 1, L0761: 1, L0772: 1, L0766: 1, L0805: 1, L0773: 1, L0749: 1, L0757: 1, L073: 1, L073: 1, H0522: 1, L073: 1, L073: 1, L0750: 1, L075: 1, L073: 1, L0753: 1, L075: 1, L073: 1, L0753: 1, L075: 1, L073: 1, L0758: 1, S0142: 1,	H0427: 1	AR089: 12, AR060: 7 H0124: 28, H0013: 8, H0547: 4, H0144: 3, L0595:
	Pro-9 to Thr-14, Ser-37 to Trp-44, Gly-79 to Thr-85, Arg-88 to Lys-139.		
	521	522	523
	250 - 666	347 - 439	238 - 300
	17	18	19
	839187	827273	847116
	HADDE71	HADD113	HADMB15
	7	×	6

## DOOS COLEDI

		·	
3, H0390: 2, S0346: 2, H0012: 2, L0565: 2, L0777: 2, S0001: 1, S0282: 1, S0442: 1, H0619: 1, S0222: 1, H0333: 1, T0039: 1, S0010: 1, S0499: 1, H0052: 1, H0546: 1, H0178: 1, H0292: 1, H0135: 1, H0591: 1, H0087: 1, H0100: 1, L0770: 1, L0531: 1, L0531: 1, L0531: 1, L0592: 1, L0651: 1, L0593: 1, L0599: 1, H0445: 1, L0592: 1, L0599: 1, H0445: 1, L0592: 1, L0599: 1 and H0352: 1.	AR060: 7, AR089: 4 L0754: 4, L0777: 2, L0755: 2, S0010: 1, H0049: 1, L0163: 1, L0771: 1, L0775: 1 and L0776: 1.	AR089: 16, AR060: 11 S0010: 1 and H0616: 1.	AR089: 8, AR060: 6 L0766: 13, L0663: 5, L0439: 3, L0747: 3, L0750: 3, H0580: 2, H0486: 2, H0013: 2, S0250: 2, L0662: 2, L0768: 2, L0527: 2, L0647: 2, L0792: 2, L0779: 2, L0596: 2, L0592: 2, L0362: 2, H0543: 2, H0556: 1, S0114: 1, H0661: 1, H0402: 1, S0420: 1, H0676: 1, H0438: 1, H0600: 1, H0497: 1, S0010: 1, L0471:
·			
	524	525	526
	171 - 236	238 - 291	146 - 313
	20	21	52
	722205	637489	823543
	HAGBQ12	HAGDW20	HAGEG10
	10	=	12

1, H0083: 1, H0267: 1, H0316: 1, H0090: 1, H0591: 1, H0038: 1, H0040: 1, L0060: 1, L0667: 1, L0373: 1, L0803: 1, L0650: 1, L0774: 1, L0775: 1, L0555: 1, L0659: 1, L0526: 1, L0529: 1, L0791: 1, L0666: 1, L0664: 1, L0666: 1, H0520: 1, H0547: 1, H0684: 1, H0520: 1, H0547: 1, H0684: 1, H0520: 1, H0547: 1, H0684: 1, L0755: 1, L0758: 1, H0445: 1, H0542: 1 and H0445: 1, H0542: 1 and	AR089: 15, AR060: 14 H0585: 12, L0439: 8, H0052: 7, H0251: 7, L0805: 7, L0776: 6, S0010: 5, L0803: 5, L0745: 5, L0809: 4, L0438: 4, L0779: 4, L0747: 3, S0222: 2, H0438: 2, T0010: 2, S6028: 2, L0455: 2, L0794: 2, L0790: 2, S0028: 2, L0742: 2, L0753: 2, S0436: 2, L0592: 2, H0650: 1, S0001: 1, S0420: 1, S0408: 1, H0013: 1, H0156: 1, T0082: 1, S0049: 1, H0263: 1, H0598: 1, H0050: 1, H0051: 1, S0049: 1, H038: 1, H0040: 1, S0386: 1, S0039: 1, L0351: 1, L0370: 1, L0770: 1, L0766: 1, L0774: 1, L0783: 1, L0788: 1, L0770: 1, L0665: 1, L0352:
	527
	515 - 550
	23
	828055
	HAGEQ79
	13

1, S0380: 1, L0740: 1, L0777: 1, L0755: 1 and L0759: 1.	AR060: 5, AR089: 3 L0438: 7, L0439: 6, L0747 4, L0005: 3, S0360: 3, H0547: 3, S0222: 2, L0105: 2, S0002: 2, S0426: 2, L0794: 2, L0659: 2, L0664: 2, L0754: 2, L0758: 2, H0506: 2, H0170: 1, H0171: 1, H0580: 1, H0455: 1, H0069: 1, H0581: 1, H0699: 1, S0003: 1, H0591: 1, L0471: 1, H0699: 1, S0422: 1, L0764: 1, L0773: 1, L0646: 1, L0773: 1, L0646: 1, L0773: 1, L0646: 1, L0773: 1, L0647: 1, S0052: 1, H0144: 1, H0682: 1, L0776: 1, L0647: 1, S0052: 1, H0144: 1, H0682: 1, H0559: 1, H0551: 1, H0555: 1, L0742: 1, H0555: 1, L0742: 1, S0434: 1 and S0452: 1	AR060: 9, AR089: 7 H0521: 5, L0777: 5, S0376: 4, H0156: 3, H0519: 3, H0436: 3, L0731: 3, H0656: 2, H0580: 2, H0036: 2, L0471: 2, H0090: 2, H0040: 2, H0551: 2, H0494: 2, S0438: 2, H0529: 2, L0809: 2, H0144: 2, S0374: 2,
	Met-1 to Lys-6.	
	528	529
	241 - 405	900 - 932
	42	25
	847120	773286
	HAGFS57	HAGHN57
	4	15

H0593: 2, H0170: 1, H0583: 1, H0650: 1, S0418: 1, S0358: 1, S0045: 1, H0619: 1, H0580: 1, H0643: 1, H0632: 1, H0648: 1, S0346: 1, H0580: 1, S0010: 1, H0590: 1, H0058: 1, H0058: 1, H0052: 1, H0615: 1, H0252: 1, H0615: 1, H0612: 1, L0455: 1, S0366: 1, H0613: 1, H0638: 1, H0634: 1, H0646:	L0790: 1, H0520: 1, H0435: 1, S0328: 1, H0539: 1, H0704: 1, S0027: 1, L0439: 1, L0750: 1, L0756: 1, L0757: 1, L0581: 1, L0595: 1, H0543: 1 and H0423: 1	AR194: 23, AR205: 21, AR206: 20, AR039: 18, AR204: 17, AR202: 17, AR204: 15, AR202: 17, AR204: 13, AR198: 12, AR310: 12, AR271: 12, AR053: 11, AR263: 11, AR033: 10, AR312: 10, AR033: 10, AR251: 10, AR309: 10, AR213: 10, AR309: 9, AR060: 9, AR309: 9, AR060: 7, AR309: 6, AR061: 7, AR249: 6, AR096: 6,
		Q
		- 237 530
		196 - 2
		26
		847013
		HAHEA15
		16

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			·
L0766: 3, H0599: 2, L0750: 2, L0753: 2, L0775: 1, L0754: 1, L0755: 1 and L0759: 1.	H0560: 1, H0561: 1 and H0542: 1.	AR060: 184, AR089: 98 H0561: 1 and L0758: 1.	AR089: 8, AR248: 7, AR265: 6, AR309: 7, AR265: 6, AR202: 6, AR212: 6, AR202: 6, AR312: 6, AR060: 5, AR194: 5, AR060: 5, AR194: 5, AR213: 4, AR052: 4, AR213: 4, AR052: 4, AR213: 3, AR310: 3, AR206: 3, AR206: 3, AR206: 3, AR206: 3, AR206: 3, AR206: 1, AR273: 2, AR204: 6, S0442: 5, S0358: 5, H0640: 4, H0640: 4, L0659: 4, H0144: 4, H0521: 4, H0171: 3, H0040: 4, L0659: 3, R055: 3, R0436: 3, L0758: 3, S0434: 3, H0551: 2, H0575: 2, H0615: 2, H0617: 2, H0616: 2, H0651: 2, H0674: 2, H0616: 2, H0651: 2, H0629: 2, L0666:
	Leu-33 to Asp-38.	Lys-89 to Glu-94.	Arg-24 to Trp-44, Leu-87 to Ser-93, Arg-119 to Trp-125, Pro-206 to Lys-211, Glu-280 to Trp-286.
	531	532	533
	192 - 308	12 - 296	605 - 1684
	27	28	53
	534670	845601	866415
	HAJAA47	HAJAY92	HAJBV67
	17	18	

2, L0740: 2, L0731: 2, L0759: 2, S0436: 2, L0362: 2, H0556: 1, S0114: 1, S0409: 1, L0002: 1, S0282: 1, S0356: 1, S0348: 1, S0408: 1, S0410: 1, H0637: 1, H0722: 1, S0046: 1, S0132: 1, S0300: 1, L0717: 1, H0411: 1, H0431: 1, H0586: 1, H0587: 1, H0036: 1, H0024: 1, H0014: 1, H0251: 1, H0596: 1, L0471: 1, H0024: 1, H0014: 1, H0375: 1, H0059: 1, L0471: 1, L0059: 1, L0655: 1, L0067: 1, H0688: 1, L0066: 1, L0369: 1, L0662: 1, L0766: 1, L0369: 1, L0662: 1, L0766: 1, L0775: 1, L0662: 1, H0683: 1, L0775: 1, L0665: 1, H0559: 1, L0665: 1, H0519: 1, S0126: 1, H0683: 1, L0775: 1, L0665: 1, H0519: 1, S0126: 1, H0683: 1, L0775: 1, L0665: 1, H0519: 1, S0126: 1, H0667: 1, L0752: 1, H0559: 1, H0667: 1, L0752: 1, L0599: 1, L0752: 1, H0569: 1, L0752: 1, H0667: 1, L0752: 1, L0599: 1, L0752: 1, L0599: 1, L0752: 1, L0599: 1, L0752: 1, L0599: 1, L0752: 1, H0667: 1, L0752: 1, L0599: 1, L0752: 1, L0599: 1, L0752: 1, S0194: 1, L0752: 1, S0194: 1, L0752: 1, R0194: 1,	H0423: 1, H0422: 1, S0384: 1, H0506: 1 and H0352: 1.		AR206: 3, AR263: 3, AR207: 3, AR312: 2,
			Asp-26 to Leu-32, Trp-62 to Asp-72,
		534	535
		284 - 400	8 - 3511
		30	31
		827275	852204
		HAJCH70	HAOAG15
		20	21

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	536	537	538
	250 - 321	262 - 273	18 - 224
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	688037	633730	839468
	HAQAI92	HAQCE11	HATBI94
	22	23	24

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			Lys-8 to Trp-13.	Val-23 to Glu-28.
	539	540	541	542
	268 - 396	296 - 409	271 - 324	93 - 221
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	631172	826098	280805	836056
	HATCB45	HATCD80	HATCI03	натен20
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2, S0003: 2, T0006: 2,	S0364: 2, H0124: 2, S0366:	2, H0135: 2, L0775: 2,	L0809: 2, L0789: 2, H0660:	2, L0753: 2, L0757: 2,	L0758: 2, L0485: 2, L0599:	2, L0601: 2, H0265: 1,	S0040: 1, H0650: 1, H0341:	1, S0212: 1, S0282: 1.	H0663: 1 H0638: 1 S0418:	000000000000000000000000000000000000000	1, 50420: 1, 50360: 1,	S0408: 1, L0149: 1, H0208:	1, S0132: 1, H0370: 1,	L0623: 1, H0013: 1, H0427:	S0280: 1. H0156: 1	H0097-1 H0575-1 H0036-	1 H0500: 1 50246: 1	10350: 1, 30340: 1,	H0318: 1, H0196: 1, H0596:	1, H0597: 1, H0231: 1,	H0009: 1, N0006: 1, L0471:	1, H0012: 1, H0024: 1,	H0373: 1, H0051: 1, H0083:	1, H0292: 1, H0428: 1,	H0604: 1, H0553: 1, H0181:	1, H0169: 1, H0163: 1,	H0090: 1, T0067: 1, H0264:	, S0038: 1, S0386: 1,	S0112: 1, L0564: 1, H0561:	, S0142: 1, S0344: 1,	L0770: 1, L0769: 1, L0637:	, L0761: 1, L0372: 1,	L0374: 1, L0521: 1, L0626:	1, L0533: 1, L0803: 1,	L0376: 1, L0806: 1, L0805:	1, L0655: 1, L0783: 1,	L0529: 1, L0666: 1, S0374:	., H0520: 1, H0547: 1,	H0519: 1, H0658: 1, S0380:
[2, S	803	2, E	T08	[2, L	1.07	2, L	800	1.S	)OH		1, 5	S04	1, S	P100		OÚA	H -	1, 17	НОЗ		00H	1, H	H03	1, H	90H		00H		S01	1, St	T01	1, L	F03	$ 1, \Gamma $	L03	1.1.	L05	1, 円	H05
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# Deemone, Deemon

1, H0521: 1, H0522: 1, H0696: 1, H0436: 1, L0609: 1, L0744: 1, L0745: 1, L0749: 1, L0777: 1, H0444: 1, L0480: 1, L0584: 1, L0595: 1, S0011: 1, H0422: 1 and H0008: 1.	AR089: 1 L0809: 4, L0766: 3, L0439: 3, H0624: 2, H0411: 2, L0794: 2, L0756: 2, L0731: 2, L0005: 1, H0599: 1, L0471: 1, S0051: 1, T0010: 1, H0266: 1, S0150: 1, L0637: 1, L0765: 1, L0803: 1, L0783: 1, H0144: 1, H0672: 1, S0392: 1, L0748: 1, L0779: 1, L0777: 1 and L0759: 1.	AR089: 8, AR060: 4 H0013: 8, L0805: 5, H0716: 4, S0010: 4, H0052: 4, H0144: 4, H0615: 3, H0547: 3, L0747: 3, H0645: 2, S0049: 2, L0776: 2, L0665: 2, H0519: 2, H0658: 2, H0660: 2, L0602: 2, H0555: 2, L0439: 2, L0750: 2, L0597: 2, H0136: 2, H0423: 2, H0624: 1, H0171: 1, H0717: 1, S0402: 1, H0294: 1, S0114: 1, S0116: 1, H0341: 1, S0212: 1, H0483: 1, H0664: 1, S0360: 1, S0046: 1, H0619: 1, H0411: 1, H0369: 1, S0222: 1, H0338: 1, H0486: 1, H0156:
1, H0521: 1, H045 H0696: 1, H045 1, L0744: 1, L07 L0749: 1, L077 1, L0480: 1, L07 L0595: 1, S001 1 and H0008: 1	AR089: 1.0809: 4.3, H0624: 2, L0094: 2, L0047: 1, H0266: L0637: 1, H0672: 1, L0779: 1,	AR089:
	543	544
	39 521 - 580	40 17 - 391
	838799	1300785
	29 HBAGD86	30 HBCJL.35

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1, H0318: 1, H0581: 1, H0046: 1, H0457: 1, H0564: 1, H0051: 1, H0416: 1, H0688: 1, H0644: 1, L0456: 1, H0135: 1, H0616: 1, H0059: 1, H0561: 1, S0344: 1, L0763: 1, L0646: 1, L0521: 1, L0766: 1, L0649: 1, L0789: 1, L0663: 1, L0438: 1, H0435: 1, S0406: 1, H0436: 1, L0612: 1, L0748: 1, L0758: 1, L0779: 1, L0758: 1, L0759: 1, L0686: 1, S0436: 1, L0759: 1, L0686: 1, S0436:		H0551: 2, L0803: 2, L0439: 2, L0750: 2, S0308: 2, L0644: 1, L0655: 1, H0479: 1, L0780: 1 and L0752: 1.	H0551: 2, L0803: 2, L0439: 2, L0750: 2, S0308: 2, L0644: 1, L0655: 1, H0479: 1, L0780: 1 and L0752: 1.	AR060: 4, AR089: 3 L0731: 20, L0747: 7, L0794: 6, L0764: 4, L0803: 4, L0759: 4, L0662: 3, L0774: 3, L0749: 3, L0756: 3, S0360: 2, H0156: 2, H0046: 2, H0181: 2, L0766: 2, L0659: 2, L0438: 2, S0126: 2, H0658: 2, L0439: 2, L0754: 2, L0777: 2,
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·	1033 - 1407	351 - 440	671 - 760	1016 - 1024
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	897937	789532	864374	691473
		HBDAB91	HBDAB91	HBGBC29
		31	32	33

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	892131	603174	846465
	HBGNC72	HBHAA05	НВНАА81
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	Arg-34 to Ser-39,  Pro-45 to Ile-55.  Loginary Exp.  A pro-45 to Ile-55.  A pro-45 to Ile-55.
	551 Arg
	1877 - 2287
	806303
	HBIAA59
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	1 88
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	HBIAC29
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		Arg-24 to Asp-31.
	553	554
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	553630	837309
	HBICW51	HBJAB02
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			Met-1 to Ala-8, Phe-42 to Asp-57,
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21		52	53
679337		909095	815649
HBJAC65		HBJBM12	HBJCR46
41		42	43

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L0803: 10, L0766: 7, S0358:	6, L0794: 6, L0775: 6,	L0809: 6, H0616: 5, L0776:	5, L0750: 5, L0591: 5,	S0408: 4, L0769: 4, L0748:	4, L0439: 4, L0747: 4,	L0731: 4, S0282: 3, L0774:	3, L0789: 3, L0666: 3,	L0438: 3, L0756: 3, L0780:	3, L0755: 3, L0757: 3,	L0759: 3, L0588: 3, H0638:	2, H0036: 2, H0590: 2,	H0318: 2, H0581: 2, H0196:	2, H0046: 2, H0154: 2,	L0163: 2, H0213: 2, S0036:	2, L0804: 2, L0375: 2,	L0655: 2, L0656: 2, L0659:	2, L0783: 2, S0374: 2,	S0126: 2, H0659: 2, H0672:	2, S0328: 2, S0152: 2,	H0521: 2, S0406: 2, L0605:	2, L0485: 2, L0362: 2,	S0026: 2, H0543: 2, H0352:	2, H0170: 1, H0171: 1,	H0713: 1, S6024: 1, H0650:	1, H0657: 1, H0656: 1,	S0116: 1, H0341: 1, S0442:	1, S0354: 1, S0360: 1,	H0580: 1, H0339: 1, S0045:	1, H0619: 1, H0437: 1,	S0222: 1, H0333: 1, H0574: [	1, L0623: 1, H0486: 1,	L0586: 1, L0021: 1, H0599:	1, H0575: 1, H0618: 1,	H0253: 1, H0230: 1, H0597:	1, H0544: 1, H0178: 1,	H0123: 1, H0050: 1, L0471:   1. H0012: 1. H0620: 1
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	1032 - 1355
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	813588
a	HBJDS79
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	559	560
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	55	56
	520401	847030
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AR053: 5, AR253: 5,	AR096: 5, AR213: 4,	AR104: 3, AR263: 2	S0440: 28, S0476: 19,	1494: 19, S0372: 16,	L0754: 16, S0132: 12,	H0046: 12, L0666: 12,	H0586: 11, S0330: 11,	S0328: 10, S0360: 9, H0587:	9. L0747: 9, H0622: 8,	S0436. 8 H0648. 7 S0356.	6 S0003: 6 H0674: 6	1,0005; 5,11007; 5,10601;	600. 3, E0302. 3, E0001.	5, 50358: 4, 50214: 4,	H0039: 4, H0031: 4, H0204:	4, L0763: 4, L0662: 4,	L0776: 4, L0777: 4, L0752:	4 S0430; 3, S0376; 3,	H0370: 3 H0600: 3 H0592:	3 H0644· 3 H0551: 3.	H0560: 3 1.0637: 3.1.0646:	3 1.0649: 3 1.0653: 3.	1.0659:3 1.0663:3 H0696:	3 \$3014: 3. \$0434: 3.	10591-3 H0662: 2, S0410:	2 H0393: 2. H0596: 2.	H0597; 2, L0483; 2, H0553;	2. H0032: 2, H0169: 2,	H0598: 2, H0090: 2, H0379:	2. L0372: 2, L0376: 2,	L0517: 2, L0783: 2, L0809:	2, 1,0665; 2, H0547; 2,	H0658: 2, H0670: 2, S0380:	2 50152.2 50406.2	2, 30122: 2, 30100: 2, \$0027: 2, 1,0744: 2, 1,0779:	2 10755-2 10599:2	S0196: 2. H0170: 1. H0171:	, H0556: 1, T0002: 1,
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1, H0231: 1, L0041: 1, H0041: 1, H0090: 1, H0123: 1, H0520: 1, H0199: 1, H0246: 1, H0199: 1, H0266: 1, H0188: 1, H0687: 1, H0288: 1, H0033: 1, H0181: 1, S0364: 1, S0366: 1, S0036: 1, H0038: 1, H0616: 1, H0264: 1, L0435: 1, T0041: 1, T0042: 1, S0448: 1, S0142: 1, S0002: 1, H0529: 1, L0796: 1, L0639: 1, L5575: 1, L5566: 1, L0761: 1, L0374: 1, L0639: 1, L0576: 1, L0648: 1, L0768: 1, L0649: 1, L0761: 1, L0555: 1, L0648: 1, L076: 1, L0648: 1, L076: 1, L0648: 1, L0776: 1, L0659: 1, L0526: 1, L0769: 1, H0520: 1, L0783: 1, H0529: 1, S0330: 1, H0539: 1, S0378: 1, S0152: 1, H0522: 1, H0694: 1, H0555: 1, S3012: 1, S0152: 1, H0544: 1, S028: 1, L0743: 1, L0779: 1, L0752: 1, H0444: 1, S0436: 1, L0581: 1, H0543: 1, H0423: 1, S0458: 1 and H0506: 1.	L0748: 8, L0749: 3, L0471: 2 and H0144: 1.	L0794: 12, H0620: 3, L0756: 2, L0759: 2, S0408:
	Ser-18 to Glu-24, Leu-121 to Asp-134, Pro-142 to Ala-154, Cys-185 to Val-203.	
	654	655
	55 - 762	212 - 448
	150	151
	834826	827796
	HE9CY05	HE9EA10
	140	141

1, S0049: 1, H0544: 1, H0012: 1, H0615: 1, H0040: 1, L0764: 1, L0803: 1, L0806: 1, L0789: 1, H0144: 1, H0547: 1, L0779: 1, L0597: 1 and L0595: 1.	AR089: 11, AR060: 9 L0748: 6, H0144: 3, S0010: 2, L0439: 2, L0749: 2, H0717: 1, H0662: 1, S6022: 1, S0222: 1, S0280: 1, L0109: 1, H0163: 1, L0639: 1, L0659: 1, L0744: 1, L0745: 1, L0747: 1, L0756: 1, L0596: 1 and S0276: 1.	AR060: 7, AR089: 4 S0418: 4, L0438: 4, L0599: 4, L0741: 3, H0581: 2, S0422: 2, L0770: 2, L0659: 2, H0520: 2, H0547: 2, L0439: 2, L0754: 2, L0747: 2, L0779: 2, S0007: 1, S0010: 1, S0049: 1, H0673: 1, H0494: 1, H0625: 1, L0769: 1, L0645: 1, L0662: 1, L0794: 1, L0766: 1, L0775: 1, L0789: 1, L0666: 1, L0775: 1, L0789: 1, L0666: 1, L0663: 1, S0374: 1, S0436: 1, L0593: 1, L0366: 1 and S0196: 1.	AR060: 4, AR089: 2 L0438: 3, T0010: 2, L0351: 2, L0748: 2, L0747: 2, S0116: 1, S0354: 1, S0007: 1, H0619: 1, H0135: 1, L0521: 1, L0774: 1, L0809: 1, H0521: 1, L0439: 1, L0755: 1, L0758: 1 and
		Val-40 to Cys-45, Lys-58 to Thr-64.	Arg-18 to Lys-26, Gly-35 to Ala-42, Gln-61 to Gly-67.
	656	657	658
	319 - 348	855 - 1064	172 - 528
	152	153	154
	633719	831464	600355
	HE9GG20	HEBCI18	HEBCY54
	142	143	144

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			Met-1 to Thr-6.
	659	099	661
	<del></del>	1211 - 1336	200 - 289
	155	156	157
	692347	840288	847064
	HEBDF77	неврQ91	HEBFR46
	145	146	147

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1, T0041: 1, S0150: 1, S0002: 1, L0761: 1, L0627: 1, L0662: 1, L0666: 1, H0547: 1, H0519: 1, H0539: 1, S0037: 1, S0206: 1, L0748: 1, L0777: 1, H0595: 1 and L0593: 1.	S0007: 1	AR060: 7, AR089: 5 H0550: 2, L0749: 2, H0483: 1, H0318: 1 and H0555: 1.	AR089: 10, AR060: 6 S0045: 1 and H0100: 1.	AR089: 15, AR060: 9 L0748: 3, S0045: 1 and L0749: 1.	AR060: 4, AR089: 3 L0750: 5, S0045: 2, S0212: 1, S0300: 1, S0010: 1, H0505: 1, S0049: 1, H0266: 1, L0598: 1, L0662: 1, L0809: 1, S0374: 1, H0696: 1 and L0758: 1.	AR060: 4, AR089: 3 L0750: 5, S0045: 2, S0212: 1, S0300: 1, S0010: 1, H0505: 1, S0049: 1, H0266: 1, L0598: 1, L0662: 1, L0809: 1, S0374: 1, H0696: 1 and L0758: 1.	AR089: 35, AR060: 16 L0717: 2, L0527: 2, S0046: 1, L0646: 1, L0748: 1, L0750: 1 and L0581: 1.	AR089: 23, AR060: 13 H0581: 2, H0457: 2 and S0116: 1.
								Ser-5 to Thr-10, Cys-36 to Glu-51.
	662	663	664	999	999	<i>1</i> 99	899	699
	106 - 234	59 - 163	215 - 277	82 - 135	147 - 215	147 - 215	440 - 472	154 - 309
	158	159	160	161	162	163	164	165
	798096	834379	693175	637624	674456	851137	834491	866171
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664 - 711	150 - 248
166	167
855935	701984
HEPBA14	HEQAH80
156	157

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3, SC	3, L(	907	90H	2, H	H05	2, H	L03(	2, L	T01	2, L	L07	2, H	90H	1. H	803	1.8	<u> </u>	1, H	H03	1, S	007	H, H	804	1, H	F01	$\begin{bmatrix} 1, \Gamma \end{bmatrix}$	T08	1,1	T08	1,1	1.05	1,1	108	1,1	THO:
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1, L0745: 1, L0747: 1, L0755: 1, H0445: 1, S0436: 1, L0592: 1, L0608: 1, L0595: 1, L0362: 1, L0361:	H0422: 1, S0424: 1 and H0352: 1.	AR060: 3, AR089: 3 H0046: 21, L0803: 4, 1,0790: 2, L0750: 2, L0777:	2, L0758; 2, L0362; 2,	3,280: 1, £0/09: 1, £0/94: 1, £0774: 1, £0809: 1 and	L0666: 1.	L0766: 9, H0521: 7,	L0731: 7, H0341: 6, L0770:	6, L0//1: 6, L0803: 6, 1 0754: 6 1 0757: 6 S0354:	5 1 0662: 5 HOS10: 5	L0439: 5, L0779: 5, L0758:	5, S0436: 5, H0009: 4,	H0673: 4, S0422: 4, L0521:	4, L0659: 4, L0438: 4,	S0028: 4, L0485: 4, L0601:	4, H0657: 3, H0638: 3,	S0418: 3, S0007: 3, S0222:	3, 30214: 3, m0329: 3, 1 0360: 3 1 0704: 3 1 0649:	3 1 0805: 3 1 0776: 3	L0809; 3, L0665; 3, H0144;	3, H0670: 3, S0406: 3,	L0756: 3, L0755: 3, L0759:	3, H0667: 3, S0420: 2,	S0358: 2, S0360: 2, H0580:	2, H0729: 2, H0733: 2,	S0476: 2, H0645: 2, S6026:	2, \$0300: 2, H0427: 2, H0156: 2, \$0010: 2, H0085:
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1, H0038: 1, H0551: 1, H0380: 1, H0623: 1, S0386: 1, T0042: 1, H0494: 1, H0561: 1, S0370: 1, H0509: 1, H0130: 1, H0641: 1, L0598: 1, L0763: 1, L0638: 1, L0796: 1, L067: 1, L0630: 1, L0373: 1, L0800: 1, L0641: 1, L0773: 1, L0661: 1, L0574: 1, L0381: 1, L0659: 1, L5623: 1, L0661: 1, L0527: 1, L0518: 1, L0651: 1, L0799: 1, L0787: 1, L0799: 1, L0787: 1, L0793: 1, S0374: 1, H0520: 1, S0126: 1, H0648: 1, H0710: 1, H0522: 1, L0749: 1, L0777: 1, L0749: 1, L0777: 1, L0393: 1, L0366: 1, S0026: 1, S0242: 1, S0276: 1, S0196: 1, H0543: 1, H0423: 1 and S0460: 1.	AR060: 7, AR089: 5 H0009: 3	AR089: 93, AR060: 55 L0747: 43, L0666: 20, L0752: 19, L0663: 18, L0439: 18, L0750: 17, L0731: 17, L0664: 13, L0665: 13, L0438: 13, L0748: 13, L0778: 13, L0662: 10, L0777: 10, L0659: 9, L0757: 9, L0775: 8, H0520: 8, H0547: 8, L0751: 8, S0436: 8, S0358:
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	692438	824057
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7	7	<u> </u>	9	<u> </u>	9	<u> </u>	5	<u>1</u>	<u>.</u>		<u> </u>		1 4	H	4		4	<u>, J</u>	4	S	<u> </u>	<u> </u>	<u> </u>	<u> </u>	<u> </u>	7	<u>6</u>	<u> </u>	2	<u> </u>	2,	<u>H</u>	2	H		H
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2, H0169: 2, H0708: 2, H0163: 2, H0087: 2, H0551:	2, H0412: 2, H0059: 2,	L0564: 2, H0625: 2, S0440:   2, L0763: 2, L0667: 2,	L0772: 2, L0773: 2, L0648:	2, L0363: 2, L0774: 2,	L0806: 2, L0633: 2, L0313:	2, LO318: 2, LO782: 2, 1 0783: 2 1 0382: 2 1 0541:	2 1.0647·2 H0690: 2.	H0670: 2, S0330: 2, H0521:	2, S0406: 2, L0741: 2,	L0743: 2, L0744: 2, L0755:	2, S0434: 2, L0588: 2,	L0590: 2, L0608: 2, L0604:	2, L0601: 2, H0668: 2,	S0196: 2, H0422: 2, H0556:	1, H0713: 1, H0717: 1,	H0716: 1, S6024: 1, S0114:	1, H0583: 1, H0650: 1,	S0116: 1, S0212: 1, S0282:	1, H0484: 1, H0402: 1,	L0005: 1, S0356: 1, S0376:	, S0444: 1, H0580: 1,	H0329: 1, H0208: 1, H0645:	1, S0278: 1, S0222: 1,	S6014: 1, H0441: 1, H0392:	1, H0587: 1, H0497: 1,	H0333: 1, H0559: 1, H0013:	1, L0021: 1, H0108: 1,	H0590: 1, H0618: 1, S0010:	1, H0318: 1, H0421: 1,	S0049: 1, H0194: 1, H0309:	, H0596: 1, H0597: 1,	H0231: 1, H0544: 1, H0546:	1, H0150: 1, H0563: 1, H0572: 1, H0570: 1, H0081:
2, H0	2, H0	L056/   2. L07	L077	2, L0	L080	2, LO	0.10	190H	2, S0	L074	2, S0	L059	2, L0	8019	1, H0	H071	1, H0	8011	1, H0	T000	1, 50	H032	1, S0	1801	1, HC	H033	1, L0	H056	1, HC	S004	1, HC	H023	1, HC H057
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H0650: 1, H0657: 1, H0638: 1, S0354: 1, S0360: 1, S0408: 1, S0278: 1, H0441: 1, H0497: 1, H0331: 1, T0109: 1, S0278: 1, H0705: 1, H0318: 1, H0687: 1, H0687: 1, H0687: 1, H0689: 1, H0655: 1, H0657: 1, L0774: 1, L0763: 1, L0774: 1, L0763: 1, L0774: 1, L0774: 1, L0523: 1, L0809: 1, L0769: 1, L0774: 1, L0523: 1, L0809: 1, L0783: 1, L0809: 1, L0783: 1, L0809: 1, L0783: 1, L0809: 1, L0783: 1, H0648: 1, H0643: 1, H0648: 1, H0643: 1, H0648: 1, H0649: 1, L0777: 1, L0749: 1, L0777: 1, L0749: 1, L0777: 1, L0769: 1, L0777: 1, L0769: 1, H0648: 1, H0648: 1, H0649: 1, L0777: 1, L0769: 1, H0649: 1, L0777: 1, L0769: 1, H0649: 1, R0849:	1 and H0506: 1. AR089: 26, AR060: 14	S0052: 1 AR089: 35, AR060: 16 S0052: 1
		Trp-31 to Arg-39, Ala-50 to Trp-57,
	805	908
	248 - 346	68 - 412
	301	302
	603910	688114
	HNGAK51	HNGAM58
	291	292

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	7, AR089:		5, AR060:	4, AK254:	3, AD271.	3, AR061:	3, AR053:	3, AR201:	3, AR096:	2, AR249:	2, AR310:	2, AK265:	2, AK253:	2, AK213:	, AK2/3: AB039:	I, AR205:	L0766: 3	2, H0620	L0754: 2	1, H0254	H0208: 1	1, H0618	H0457: 1	1, H0271	H0063: 1	1, T0042	L0761: 1,	, L0805: 1, L0655: 1,	S0188: 1	I allu IIO	
	AR060: 7 S0052: 1			AR052: 4									AR19/: 2	7 -	AK308: 1 AP194: 1		4	2, H0402: 2, H0620:	H0024: 2, L0754: 2, H0656:	1, H0484: 1, H0254: 1,	S0360: 1, H0208: 1, H0393:	1, S0222: 1, H0618: 1	H0194: 1, H0457: 1, H0123:	1, H0051: 1, H0271: 1	H0182: 1, H0063: 1, H0087:	1, L0351: 1, T0042: 1	S0448: 1, L0761: 1, L0378:	, L0805:	H0539: 1, S0188: 1, S0146:	., no.43.	
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eu-93, Gly-113.	ilu-24.	rg-33.	er-39,	iln-61, Ser 120	Jel-139.																									6	er-39, 31n-61, Glv-130
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	807	808	608																											0,0,	1017
	- 187	- 384	237 - 965																			•								1	- 629
	47	205	237												<u> </u>					••										-	23]
	303	304	305																											1	513
	532614	825389	1145071							•																					866177
	HNGBH53	HNGD038	HNGDX18																												
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	His-26 to Ser-32.		Ala-8 to Gly-20.							Ser-41 to Ser-48, Arg-61 to Trp-68.		
810	811	812	813	814	815	816	817	818	819	820	821	822
73 - 126	58 - 192	30 - 98	184 - 282	181 - 387	25 - 54	221 - 247	252 - 473	415 - 552	452 - 583	611 - 835	467 - 571	544 - 621
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566863	815678	535723	638116	597449	88668	561572	852178	836061	838613	834949	836063	834942
HNGDY34	HNGEA34	HNGEQ75	HNGGA68	HNGGP65	HNGHZ69	HNGIV64	HNGJB41	HNGKT41	HNGMW45	HNGNK44	HNGNO53	HNGPJ25
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			Pro-40 to Tyr-46.	Pro-23 to Gln-34.	
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78 - 131	298 - 663	239 - 433	215 - 394	192 - 317	91 - 111
319	320	321	322	323	324
836157	834487	597451	634589	778392	855957
HNHEN82	HNHFE71	HNHGK22	HNHHB10	HNHKS19	HNTBT17
309	310	311	312	313	314

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	835084	868327	792928
	HOABP31	HOABP31	HOACG07
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			H0585: 5, H0677: 5, H0427:	
			4, L0749: 4, L0731: 4,	
			H0618: 3, H0617: 3, L0769:	
			3, L0800: 3, H0556: 2,	
			H0141: 2, H0716: 2, H0587:	
			2, S0049: 2, H0052: 2,	
			H0123: 2, H0266: 2, H0135:	
			2, H0412: 2, S0142: 2,	
			L0761: 2, L0794: 2, L0649:	
			2, L0657: 2, L0659: 2,	
			L0663: 2, L0665: 2, H0689:	
			2, H0506: 2, H0713: 1,	
_			H0657: 1, H0483: 1, H0255:	
			1, H0661: 1, H0638: 1,	 
			S0356: 1, S0442: 1, S0360:	
			1, H0580: 1, H0722: 1,	
			S0046: 1, S0278: 1, H0441:	
			1, H0438: 1, H0559: 1,	
			H0013: 1, H0253: 1, H0546:	
			1, H0545: 1, H0041: 1,	
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			1, S0051: 1, H0292: 1,	
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	329	330	331	332	333
	655356	520196	825283	836069	789396
	HODAG07	HODBB70	HODB V05	HODCZ32	НОЕВК60
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	838	839	840	841	842
	48 - 263	138 - 257	230 - 385	309 - 371	21 - 140
	334	335	336	337	338
	836646	762821	897611	772573	638293
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	843	844	845	846
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	339	340	341	342
	839270	862049	688055	562778
	HORBV76	HOSDO75	HOSEC25	HOSEI81
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H0539:	AR08	H003	6. S01	H065	4, L0	H062	3, L05	H053	3. LO	H017	2, H0	S0046	2, H00	H059	2, H0	S0210	2, L0	T055	2, LO	)SE033(	2, L0	H039	1, 50	T000;	1, 50	S047(	1, H0	H057	1, HO	H025	1, H0	F016.	1, S02	H064	1, H0674 H0551: 1
	7																																		
	848 - 934 847				•			-					-																				-		· · · · ·
	343 84																											-							
	795132																																		
	HOSEJ94																																		
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1, T0042: 1, H0494: 1, H0561: 1, S0438: 1, S0422: 1, UNKWN: 1, L0520: 1, L0625: 1, L0637: 1, L0627: 1, L0772: 1, L0764: 1, L0773: 1, L0521: 1, L0662: 1, L0768: 1, L0522: 1, L0650: 1, L0790: 1, L4501: 1, L0666: 1, L0663: 1, L0664: 1, S0216: 1, S0374: 1, H0659: 1, H0660: 1, H0672: 1, S0328: 1, H0696: 1, H0478: 1, S3012: 1, S0027: 1, S0028: 1, L0750: 1, L0596: 1, L0592: 1, L0596: 1, L0592: 1, L0608: 1, L0361: 1, S0192: 1, H0423: 1 and L0697: 1.	S0040: 1, T0042: 1 and S0292: 1.	H0052: 17, L0745: 11, L0748: 10, H0547: 7, L0439: 7, L0755: 6, L0771: 5, L0774: 5, L0662: 4, L0746: 4, L0777: 4, L0163: 3, H0059: 3, H0100: 3, L0775: 3, L0741: 3, H0261: 2, H0333: 2, H0194: 2, H0545: 2, H0012: 2, H0617: 2, H0135: 2, L0770: 2, L0665: 2, L0438: 2, H0520: 2, L0747: 2, L0752: 2, L0753: 2, S0040: 1, L0717: 1, H0437: 1, H0550: 1, S6016: 1, H0497: 1, H0574:
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	200 - 301	70 - 336
	344	345
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1, T0114: 1, H0590: 1, H0004: 1, S0010: 1, S0346: 1, T0048: 1, S0049: 1, H0052: 1, H0263: 1, H0596: 1, H0373: 1, H0244: 1, H0373: 1, T0010: 1, H0373: 1, H0373: 1, H0373: 1, H0373: 1, H0169: 1, H028: 1, H0169: 1, H0264: 1, H0379: 1, H0616: 1, H0598: 1, H0616: 1, H0698: 1, H0591: 1, H0616: 1, H0628: 1, H0626: 1, H0626: 1, H0626: 1, H0626: 1, H0649: 1, L0370: 1, H0640: 1, L0371: 1, L0649: 1, L0649: 1, L0649: 1, L0649: 1, L0648: 1, H0679: 1, L0649: 1, L0648: 1, H0679: 1, L0649: 1, L0648: 1, H0672: 1, H0648: 1, H0673: 1, L0774: 1, L0774: 1, L0774: 1, L0774: 1, H0648: 1, H0672: 1, H0648: 1, H0672: 1, H0648: 1, H0673: 1, L0774: 1, L0775: 1, H0648: 1, H0675: 1, L0776: 1, H0666: 1, S0406: 1, H0655: 1, L0776: 1, L0776: 1, H0696: 1, S0406: 1, H0655: 1, L0776: 1, L0776: 1, H0696: 1, S0406: 1, H0655: 1, L0776: 1, H0696: 1, S0406: 1, H0655: 1, L0776: 1, L0776: 1, H0696: 1, S0406: 1, H0655: 1, L0776: 1, L0776: 1, L0776: 1, H0696: 1, S0406: 1, H0655: 1, L0776: 1, L0776: 1, L0776: 1, L0776: 1, H0696: 1, S0406: 1, H0655: 1, L0776: 1, L0776: 1, L0776: 1, L0776: 1, H0696: 1, S0406: 1, H0655: 1, L0776: 1, L077	1, L0780: 1, L0753: 1, H0595: 1, S0436: 1, L0592: 1, S0011: 1, S0026: 1, H0653: 1, S0276: 1 and	H0352: 1. AR060: 15, AR089: 14
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	420	421 63
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	HT5GR59	HTAEI78
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422	423
566861	812332
HTDAA78	HTEAG62
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	HTEDF18	HTEDI28
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L0483: 1, H0553: 1, H0674: 1, H0038: 1, L0564: 1, S0382: 1, H0538: 1, L0763: 1, L0763: 1, L0763: 1, L0638: 1, L0765: 1, L0771: 1, L0649: 1, L0522: 1, L0775: 1, L0375: 1, L0659: 1, L0438: 1, H0690: 1, H0648: 1, S0330: 1, H0521: 1, S0032: 1, L0756: 1, L0753: 1, L0755: 1, L0753: 1, L0755: 1, L0753: 1, H0423: 1, H0422: 1, S0462: 1, and S0460: 1.	H0253: 4, L0779: 2, H0618: 1, H0050: 1, H0038: 1, L0151: 1, L0758: 1 and H0445: 1.		AR060: 5, AR089: 2 H0038: 5, L0758: 3, L0770: 2, H0539: 2, L0731: 2, T0049: 1, S0358: 1, H0574: 1, H0012: 1, H0428: 1, H0135: 1, L0764: 1, L0522: 1, L0803: 1, L0650:
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	838621	762846	753425
	HTEDS12	HTEED26	HTEED26
	418	419	420

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	433				434						435																		436					
	764835				827700						862066													_					806461					
	HTEEW69				HTEGS07						HTEGS11																		HTEHA56					
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201 1,1 201,	H	5,1 1,0 1,0 1,0 1,0 1,0 1,0 1,0
	942	943
	101 - 172	171 - 287
	438	439
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)HO:	4, I	803	3, I	)OH	2, 1	)OH	2.1	01		7,7	101 101	2,1					217		1,1	OH	1,1	OH	1, 8	H0.	1,1	803	1,1	HO	1,1	0H	1,1		1,1		1,1	TO	1,1	HO	1,8
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	954 Arg-29 to														,								-											
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	Met-1 to Arg-12, Thr-19 to Leu-27, Asp-72 to Val-79, Arg-89 to Pro-94, Lys-102 to Ser-111, Glu-116 to Arg-122, Lys-134 to Pro-142, Ser-146 to Ser-151, Gly-177 to Asp-196.					
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	205 - 825	91 - 120	209 - 304		340 - 411	1802 - 1915
	451	514	452		453	454
	723799	566786	535937		846063	835069
	HTLEV48		HTLFA13		HTLFI73	HTLG189
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				H0413: 2, H0100: 2, H0494:		
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				2, L0595: 2, H0352: 2,		
				S0040: 1, L0760: 1, S0116:		
				1, S0282: 1, H0638: 1,		
				S0418: 1, S0356: 1, H0393:		•
				1. H0549; 1. H0550; 1.		
				H0333: 1. H0486: 1. T0114:		
				1 H0250-1 H0069-1		
				S0280: 1 H0156: 1 H0599:		
				1 H0575· 1 H0036· 1		
				T, 1102/3: 1, 110333: 1,		
				1 NOOOK: 1, HOSE3: 1, HOLVE:	-	
				HO107: 1 HO100: 1 HO051:		
				1 100001, 1, 110199, 1, 110001.		
				1, HUUOU: 1, HUI 66: 1, HADOO: 1 HAD 84: 1 HO478:		
		•		1 ED623: 1 1 D483: 1		
		•		1, F10022: 1, L0463: 1,		
				H0124: 1, H0135: 1, H0163:		
				1, H0040: 1, H0264: 1,		
				H0412: 1, L0564: 1, H0130:		
				1, H0641: 1, S0144: 1,		
				S0002: 1, L0763: 1, L0761:		
				1, L0372: 1, L0643: 1,		
•				L0764: 1, L0771: 1, L0648:		
				1, L0767: 1, L0803: 1,		
				L0804: 1, L0375: 1, L0378:		
	•			1, L0659: 1, L0544: 1,		
				L0665: 1, L0352: 1, H0670:		-
				1, S0328: 1, S0330: 1,		
				S0152: 1, H0134: 1, H0436:		
				1, S0027: 1, L0747: 1,		
				L0756: 1, L0777: 1, L0753:		
				1, S0260: 1, H0445: 1,		

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L0597: 1, H0653: 1 and S0194: 1.	H0253: 7, H0618: 4, H0620: 3, L0794: 3, L0769: 2, L0768: 2, L0439: 2, H0327: 1, H0051: 1, S0250: 1, S0036: 1, L0639: 1, L0761: 1, L0635: 1, L0791: 1, L0664: 1, L0438: 1, H0539: 1, L0741: 1, L0747: 1, L0750: 1, L0756: 1 and L0753: 1.	AR089: 40, AR060: 25 H0616: 14, H0038: 12, H0618: 7, H0253: 5, L0758: 5, L0768: 4, H0411: 2, L0779: 2, L0151: 1, L0697: 1 and S0398: 1.	AR089: 40, AR060: 25 H0616: 14, H0038: 12, H0618: 7, H0253: 5, L0758: 5, L0768: 4, H0411: 2, L0779: 2, L0151: 1, L0697: 1 and S0398: 1.	AR089: 40, AR060: 25 H0616: 14, H0038: 12, H0618: 7, H0253: 5, L0758: 5, L0768: 4, H0411: 2, L0779: 2, L0151: 1, L0697: 1 and S0398: 1.	AR089: 40, AR060: 25 H0616: 14, H0038: 12, H0618: 7, H0253: 5, L0758: 5, L0768: 4, H0411: 2, L0779: 2, L0151: 1, L0697: 1 and S0398: 1.	AR089: 40, AR060: 25 H0616: 14, H0038: 12,
	Pro-4 to Gly-9.	Phe-30 to Lys-37, Pro-43 to Lys-75.	Phe-30 to Lys-37, Pro-43 to Lys-75.			
	959	096	961	962	963	964
	933 - 1049	642 - 869	644 - 871	644 - 871	644 - 871	644 - 871
	455	456	457	458	459	460
	843506	834946	842691	870167	886780	891533
	HTLF11	HTLF12	HTLF12	HTLF12	HTLE12	HTLIF12
	445	446	447	448	449	450

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	Phe-30 to Lys-37, Pro-43 to Lys-75.					Tyr-21 to Phe-26, Glu-58 to Trp-66.
	596	996	296	896	696	970
	644 - 871	193 - 285	534 - 599	61 - 144	89 - 193	228 - 443
	461	462	463	464	465	466
	901225	266880	831967	638623	664508	823126
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	Ile-35 to Cys-42.	7 7 7	Gly-26 to Val-32.	Gly-26 to Val-32.	
	974	975	976	77.6	978
	439 - 630	198 - 362	134 - 310	221 - 397	52 - 153
	470	471	472	473	474
	732808	853621	806212	762851	840596
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	HTSFJ32	HTTCB60
	465	466

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1 and H0422: 1.	AR089: 168, AR060: 110	L0666: 19, L0664: 19,	S0360: 17, S0344: 17,	LU648: 17, SU358: 16,	L0053: 14, L0390: 13,	1.0663: 10.1.0740: 0.1.0775:	8 1 0599: 8 S0376: 7	H0046: 7. H0486: 6. H0597:	6, S0126: 6, L0439: 6,	L0752: 6, S0116: 5, S0140:	5, H0581: 5, S0328: 5,	L0748: 5, H0543: 5, H0423:	5, H0657: 4, S0212: 4,	H0617: 4, H0087: 4, S0372:	4, L0374: 4, L0651: 4,	H0555: 4, L0744: 4, L0754:	4, L0747: 4, T0049: 3,	S0278: 3, H0031: 3, H0641:	3, S0144: 3, L0646: 3,	L0375: 3, L0776: 3, L0606:	3, L0661: 3, L0657: 3,	S0428: 3, H0518: 3, H0521:	3, S3014: 3, L0742: 3,	L0743: 3, L0750: 3, L0753:	3, L0362: 3, L0601: 3,	S0026: 3, H0265: 2, H0556:	2, T0002: 2, H0686: 2,	S0114: 2, H0402: 2, S0410:	2, S0300: 2, T0060: 2,	H0575: 2, H0274: 2, H0318:	2, H0085: 2, H0231: 2,	H0083: 2, H0271: 2, H0188:	2, H0688: 2, H0553: 2,	H0068: 2, H0509: 2, S0142:	2, S0002: 2, L0369: 2,
	286																																		
	65 - 520										•	•									-														
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	Prò-70 to Ser-89, Ser-92 to Ser-115.	
	8886	686
	49 - 447	216 - 416
	484	485
	853408	658730
	HTXDD61	HTXDG92
	474	475

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	138:		139:		)52:		-90		341:		-69		41.	 -	- 199		7.	<del></del>	<u>-</u>		Ç	4 		 .: 0		 		:981		327:		115:		:98		:63:	
~	H0618: 7, L0758: 7, L0438:	. 5,	H0556: 4, H0059: 4, L0439:	.; ,	S0410: 3, H0253: 3, H0052:	7:3,	L0764: 3, L0768: 3, L0806:	 ش	L0731: 3, H0583: 2, H0341	, <b>6</b>	S0222: 2, H0013: 2, H0069:	2	H0087-2 1,0351-2 T0041-	, ,	L0773: 2. L0662: 2. L0766:	; ; ;	z, zococy. z, zococy. z, I 0793· 2 I 0665· 2 H0547·	, 116	1. 2,	LO/42: 2, LO/46: 2, FIO342: 2 SOOAO: 1 HO717: 1	1,	HU/16: 1, SU114: 1, 10049:	٠. L,	HU381: 1, HU863: 1, SU36U:	: I,	S0132: 1, S0300: 1, S0278:	.; ;	H0431: 1, H0392: 1, H0486:	<del>_</del>	H0581: 1, H0374: 1, H0327:	: <b>T</b> :	H0012: 1, H0024: 1, H0015:	_ <u>_</u> ;	H0188: 1, H0292: 1, H0286:	<u>-</u> :	H0181: 1, H0135: 1, H0163:	1, H0040: 1, H0063: 1,
1.0777-11 1.0754-8	758:7	6, H0144: 5, L0747: 5	059: 4	4, L0601: 4, H0265: 3	53:3	3, H0620: 3, H0617: 3,	68: 3	3, L0744: 3, L0749: 3.	383: 2	2, S0358: 2, S0046: 2,	113:2	2, S0049; 2, H0150; 2	\$51.2	2. H0529; 2. L0771; 2	62.2	2 1.0803 2 1.0809 3	65.0	2 UOS 10: 2 UOG 50: 2, II	10001 2001	2 S0040: 1 H0717:	7 7 7	14: L	1, H003 /: 1, H0036: 3	703: 1	I, H0/22: I, S0007: I	90: 1,	1, H0261: 1, H0550: 1	92: 1	1, T0114: 1, S0010: 1	374: 1	1, H0545: 1, H0457: 1	24: 1	1, H0510: 1, H0594: 1	92: 1	l, H0622: 1, H0553: 1,	35: 1	1, H0040: 1, H0063: 1,
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7777	618:	<b>4014</b>	256: 4	.0601	110:3	10620	764: 3	.0744	/31:3	30358	22:2	0049	787.	10525	73: 2	0803	703.7	7.5. Z	27.57	7.74	1.47	10: 1	10007	10101	10/22 20/22	32: 1	10261	31: 1	0114	381: 1	10545	112: 1	01501	88: 1	[0622	81: 1	00040
1	HOH	6, F	HO	4, I	S04	3, F	<u>[0</u>	3, I	<u> </u>	2, 5	S02	2.5	HO	2. F	$\frac{1}{10}$	2 .	1,5	) ;	7, 1	3 6	; ;	) -	1, 1		I, E	<u>S</u>	], H	H 6	I, T	H05	1, H	HOC	1, H	H 101	1, H	H01	1, H
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				Met-1 to Pro-6, Gly-73 to Thr-78.
	066	991	992	993
	178 - 267	192 - 281	108 - 158	330 - 566
	486	487	488	489
	581521	853410	637774	834438
	HTXET11	HTXFA72	HTXJY08	HTXKF95
	476	477	478	479

L0754: 41, L0747: 8, L0755: 5, L0659: 4, H0265: 2, H0556: 2, H0586: 2, L0471: 2, H0553: 2, L0764: 2, L0662: 2, L0794: 2, L0748: 2, L0751: 2, L0749: 2, L0750: 2, H0305: 1, S0358: 1, S0046: 1, H0441: 1, H0599: 1, H0569: 1, H0050: 1, H0616: 1, L0770: 1, L0769: 1, L0800: 1, L0644: 1, L0363: 1, L0806: 1, L0783: 1, L0666: 1, L0665: 1, H0144: 1, H0555: 1, S3012: 1, L0779: 1, L0731: 1, L0605: 1, L0599: 1, L0603: 1, H0543: 1, H0422: 1 and H0566: 1	AR060: 7, AR089: 4 L0439: 6, H0556: 2, S0007: 2, L0744: 2, L0740: 2, L0731: 2, S0442: 1, L0021: 1, H0618: 1, H0253: 1, H0041: 1, L0770: 1, L0800: 1, L0766: 1, L0803: 1, L0375: 1, L0807: 1, L0382: 1, L0791: 1, L0793: 1, L0352: 1, S0432: 1, L0741: 1 and L0779: 1.	AR060: 26, AR089: 7 L0764: 5, L0771: 5, H0506: 4, L0374: 3, S0434: 3, S0356: 1, S0408: 1, H0264: 1, L0372: 1, L0783: 1, L0532: 1 and L0663: 1.
	Pro-19 to Ser-28.	
	994	995
	319 - 432	287 - 367
	490	491
	834881	801938
	HTXMZ07	HUFCL31
	480	481

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AR089: 14, AR060: 9	H0052: 13, S0360: 8,	L0748: 8, H0619: 6, L0659:	6, L0665: 6, L0759: 6,	L0789: 5, L0743: 5, L0752:	5, S0346: 4, H0059: 4,	L0662: 4, L0805: 4, H0521:	4, L0717: 3, H0599: 3,	H0644: 3, L0761: 3, L0776:	3, S0028; 3, L0744; 3,	L0754: 3, L0749: 3, L0731:	3, L0757; 3, S0001; 2.	S0354: 2, H0261: 2, H0586:	2, S0010: 2, H0620: 2.	L0771: 2, L0804: 2, L0774:	2, L0806: 2, L0809: 2.	L0664: 2, H0547: 2, H0539	2, H0555; 2, L0747; 2,	L0750: 2, L0758: 2, S0434:	2, L0596: 2, L0604: 2,	H0171: 1, S0040: 1, H0713:	1, H0656: 1, S0212: 1,	L0005: 1, S0356: 1, H0728:	1, H0733: 1, S0046: 1,	S0278: 1, H0370: 1, H0392:	1, H0602: 1, H0592: 1,	H0574: 1, H0013: 1, S0280:	1, H0575: 1, T0082: 1,	H0581: 1, H0544: 1, H0046:	1, H0009: 1, H0081: 1,	H0051: 1, H0266: 1, H0179:	1, H0290: 1, H0286: 1,	S0250: 1, S0366: 1, S0036:	1, H0135: 1, H0591: 1,	H0038: 1, H0551: 1, H0264:	1, H0488: 1, T0004: 1,	H0100: 1, H0429: 1, H0334:	1, H0386: 1, S0144: 1,
Ser-32 to Arg-39.	•																																				
966																																					
273 - 392												•																					1				
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S0344: 1, S0002: 1, L0763: 1, L0667: 1, L0764: 1, L0773: 1, L0794: 1, L0766: 1, L0803: 1, L0793: 1, L0666: 1, S0053: 1, H0144: 1, L0352: 1, H0520: 1, H0660: 1, H0672: 1, S0328: 1, H0696: 1, S0404: 1, S0406: 1, H0436: 1, S0390: 1, H0436: 1, S0390: 1, L0777: 1, S0031: 1, S0050: 1, L0777: 1, S0031: 1, S0260: 1, L0584: 1, L0591: 1 and H0506: 1.	AR060: 6, AR089: 3 H0266: 1 and H0059: 1.	AR089: 14, AR060: 8 S0053: 4, H0673: 3, H0618: 2, H0179: 2, H0674: 2, S0216: 2, H0521: 2, S0031: 2, H0556: 1, S0116: 1, H0305: 1, H0619: 1, H0550: 1, H0069: 1, H0635: 1, H0318: 1, H0309: 1, H0083: 1, H0271: 1, H0090: 1, H0634: 1, H0059: 1, S0002: 1, S0052: 1, S0428: 1, H0144: 1, S0152: 1 and L0740: 1.	AR245: 5, AR194: 4, AR061: 3, AR251: 3, AR201: 2, AR205: 2, AR198: 2, AR039: 2, AR055: 2, AR250: 2, AR204: 2, AR060: 1, AR312: 1, AR311: 1, AR243: 1, AR186: 1, AR089: 1, AR263: 1
			Phe-166 to Arg-174, Ser-191 to Tyr-196.
	266	866	666
	214 - 315	187 - 285	74 - 661
	493	494	495
	566823	570896	894699
	HUKDF20	HUKDY82	HUSCJ14
	483	48 48	485

L0777: 8, L0766: 7, L0741:	L0754: 5, L0744: 4, L0757:	4, S0192: 4, H0677: 4,	H0556: 3, S0360: 3, S0410:	3, H0013: 3, H0052: 3,	L0769: 3, L0775: 3, L0776:	3, L0756: 3, L0752: 3,	L0604: 3, H0265: 2, S0040:	2, H0599: 2, H0545: 2,	H0266: 2, H0030: 2, H0617:	2. H0135: 2. L0771: 2.	L0662: 2, L0806: 2, L0805:	2, L0659: 2, L0666: 2,	L0665: 2, H0520: 2, H0547:	2, H0519: 2, H0659: 2,	S0404: 2, L0743: 2, L0758:	2, L0596: 2, L0605: 2,	L0485: 2, H0171: 1, H0713:	1, S0134: 1, S0218: 1,	H0657: 1, H0656: 1, S0212:	1, H0663: 1, S0420: 1,	S0408: 1, S0132: 1, S0476:	1, H0393: 1, H0587: 1,	T0040: 1, T0060: 1, H0575:	1, H0309: 1, H0009: 1,	L0471: 1, H0620: 1, H0510:	1, H0290: 1, S0250: 1,	S0022: 1, T0023: 1, L0055:	1, H0634: 1, H0488: 1,	H0268: 1, T0041: 1, T0042:	1, H0538: 1, S0210: 1,	L0763: 1, L0639: 1, L0764:	1, L0794: 1, L0649: 1,	L0804: 1, L0650: 1, L0774:	1, L0809: 1, L0793: 1,	L0664: 1, H0144: 1, H0593:	1, S0122: 1, H0435: 1,
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H0521: 1, S0406: 1, H0555: 1, L0740: 1, L0747: 1, 1, L0759: 1, L0779: 1, L0731: 1, L0759: 1, S0031: 1, S0434: 1, S0436: 1, L0601: 1, S0106: 1, H0665: 1, H0667: 1 and S0276: 1.	AR252: 82, AR250: 77, AR253: 70, AR254: 37, AR309: 22, AR264: 17, AR308: 16, AR312: 15, AR263: 15, AR096: 13, AR211: 11, AR271: 10, AR213: 8, AR243: 7, AR245: 7, AR053: 7, AR246: 6, AR272: 6, AR089: 6, AR212: 6, AR099: 6, AR212: 6, AR099: 6, AR212: 6, AR099: 3, AR039: 3, AR204: 4, AR033: 4, AR204: 4, AR033: 4, AR204: 4, AR039: 3, AR205: 2, AR104: 3, AR055: 2 L0766: 4, S0358: 3, H0266: 3, S0356: 2, S0045: 2, L0777: 2, L0731: 2, H0422: 2, H0422: 2, H0616: 2, L0794: 2, H0421: 1, H0491: 1, H0486: 1, H0377: 1, H0491: 1, H0378: 1, B0008: 1, H0378: 1, B0008: 1, H0378: 1, B0008: 1, H0378: 1, B0008: 1, H0328: 1, B0006: 1, H0644: 1, H0032:
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	486

1, H0090: 1, H0038: 1, H0087: 1, H0264: 1, H0268: 1, H0412: 1, S0422: 1, H0529: 1, L0521: 1, L0803: 1, L0659: 1, L0666: 1, H0710: 1, H0518: 1, H0521: 1, S0176: 1, S0406: 1, S3014: 1, L0439: 1, L0758: 1 and H0543: 1.	AR089: 34, AR060: 23	L0748: 4, H0622: 3, L0777: 3, H0624: 2, H0013: 2, H0520: 2, H0539: 2, L0439: 2, L0754: 2, L0747: 2, L0757: 2, L0758: 2, L0593: 2, L00002: 1, H0664: 1, H0580: 1, S0007: 1, H0497: 1, H0333: 1, H0599: 1, H0581: 1, L0483: 1, H0598: 1, H0040: 1, H0412: 1, L0351: 1, T0041: 1, L0769: 1, L0771: 1, L0662: 1, L0766: 1, L0381: 1, L0866: 1, L0766: 1, L0659: 1, L0766: 1, L0659: 1, L0766: 1, L0659: 1, L0760: 1, L0749: 1, L0750: 1, L0779: 1, L0752: 1, L0480: 1, L0591: 1 and H0543: 1.	AR060: 5, AR089: 3 H0393: 1, H0056: 1 and L0662: 1.	AR089: 4, AR060: 2 H0547: 12, L0794: 10, H0251: 9, L0439: 8, L0731:
	Arg-21 to Ser-27, Ile-36 to Asp-41.			
	1001	1002	1003	1004
	500 - 640	83 - 151	196 - 213	223 - 312
	497	498	499	500
·	684975	762858	564853	830432
	HUSGU40	HUSIR18	HUVDJ48	HWAAI12
	487	88	489	490

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		H0351: 5, L0750: 5, S0356:	
		4, L0768: 4, L0766: 4,	
		L0805: 4, L0809: 4, L0777:	
		4, L0758: 4, L0596: 4,	
		S0410: 3, H0013: 3, H0009:	
		3, H0594: 3, T0006: 3,	
	_	H0124: 3, T0041: 3, H0529:	
		3, L0769: 3, L0666: 3,	
		H0144: 3, H0520: 3, L0749:	
		3, H0661: 2, H0305: 2.	
		S0360; 2, 1,0103; 2, H0581;	
		2, S0049; 2, H0052; 2.	
	_	1.0157: 2.1.0471: 2. H0031:	
		2. H0087: 2. H0100: 2.	
		80344: 2, 1,0763: 2, 1,0761:	
		2, 1,0662; 2, 1,0803; 2,	
		L0806: 2, L0664: 2, H0436:	
		2. S0028: 2. 1.0742: 2	
		L0756: 2, L0752: 2, L0605:	
		2, L0595; 2, H0543; 2,	
		H0556: 1, T0002: 1, S0040:	
		1, H0717: 1, H0716: 1,	
		H0294: 1, S0134: 1, H0341:	
		1, H0402: 1, S0354: 1,	
		L0717: 1, S0278: 1, L0394:	
-		1, H0549: 1, S0222: 1,	
		H0333: 1, L0622: 1, H0486:	
		1, H0250: 1, H0635: 1,	
		H0575: 1, S0010: 1, H0421:	
		1, H0085: 1, H0597: 1,	
		H0545: 1, H0566: 1, H0620:	
		1, H0271: 1, H0687: 1,	
		H0615: 1, H0622: 1, H0673:	
		1, H0674: 1, H0412: 1,	
		H0413: 1, H0056: 1, T0042:	
		1, H0130: 1, H0646: 1,	
		S0144: 1, S0208: 1, L0770:	

			•
1, L0796: 1, L0667: 1, L0772: 1, L0373: 1, L0372: 1, L0800: 1, L0645: 1, L0764: 1, L0648: 1, L0767: 1, L0650: 1, L0657: 1, L0517: 1, L0789: 1, L0790: 1, L0665: 1, H0690: 1, H0658: 1, H0670: 1, H0672: 1, S0378: 1, S0380: 1, H0521: 1, S3012: 1, S0390: 1, S0027: 1, L0743: 1, L0779: 1, L0755: 1, L0759: 1, S0031: 1, S0436: 1, L0601: 1, S0276: 1, H0542: 1 and S0424: 1.	AR060: 2, AR089: 1 L0717: 2, H0580: 1, S0222: 1, L0662: 1, H0436: 1, L0748: 1, H0445: 1 and S0308: 1.	AR060: 1, AR089: 1 H0580: 1	AR089: 11, AR060: 5 H0635: 7, L0794: 6, H0556: 4, S0414: 4, H0521: 4, H0634: 3, L0779: 3, H0265: 2, S0134: 2, S0360: 2, H0619: 2, H0069: 2, H0575: 2, H0688: 2, H0056: 2, S0002: 2, L0665: 2, S0216: 2, H0519: 2, L0751: 2, L0758: 2, L0593: 2, H0422: 2, S0114: 1, S0116: 1, H0300: 1, S0356: 1, H0580: 1, S0045: 1, S0046: 1, H0643: 1, H0250: 1, H0581: 1, S0049: 1, L0045: 1, H0622: 1, H0031: 1,
	Ala-21 to Ser-31.	Lys-45 to Pro-51, Arg-80 to Arg-85.	Ser-30 to Gly-36.
	1005	1006	1007
	222 - 353	378 - 650	253 - 405
	501	502	503
	689121	722259	762860
	нwввQ70	HWBCN36	HWBDJ08
	491	492	493

H0644: 1, H0551: 1, H0264: 1, H0623: 1, H0641: 1, H0646: 1, L0763: 1, L0536: 1, L0766: 1, L0653: 1, L0555: 1, H0134: 1, L0777: 1, L0755: 1, H0542: 1, H0543: 1 and H0423: 1.	AR060: 184, AR089: 165 S0114: 1 and H0580: 1.	AR089: 61, AR060: 49, AR198: 5, AR194: 4, AR096: 3, AR310: 3, AR265: 3, AR213: 2, AR312: 2, AR249: 2, AR312: 2, AR063: 2, AR052: 2, AR104: 2, AR205: 1, AR039: 1 H0580: 1, S0300: 1, H0600: 1, L0783: 1, L0438: 1, L0439: 1 and L0758: 1.	AR060: 28, AR089: 14 H0556: 19, H0265: 15, S0418: 10, S0358: 9, S0440: 9, L0755: 9, S0420: 8, L0752: 7, H0253: 6, L0751: 6, L0747: 6, L0750: 6, L0596: 6, S0212: 5, H0618: 5, H0545: 5, H0012: 5, H0617: 5, H0413: 5, L0740: 5, L0601: 5, H0295: 4, S0360: 4, H0039: 4, H0494: 4, H0641: 4, L0764: 4, L0776: 4, S0406: 4, L0758: 4, H0445: 4, H0657: 3, H0483: 3, S0356: 3, S0376: 3, S0408: 3, S0346: 3, H0040: 3, S0344: 3, L0637: 3, H0547: 3, H0658: 3,
	1008	1009	1010
	267 - 278	242 - 349	866 - 964
	504	505	506
	827312	821335	796743
	HWBFX16	HWDAC26	HWDAG96
	494	495	496

-			110660 2 80208 2 110623	
			HU00U: 3, 3U326: 3, HU322:	
			3, L0743: 3, L0749: 3,	
			L0756: 3, L0731: 3, L0757:	
			3, S0040: 2, H0713: 2.	
			H0294: 2. H0341: 2. H0484:	
			2, H0661: 2, H0305: 2,	
			H0125: 2, H0580: 2, H0586:	
			2, H0587: 2, H0052: 2,	
			H0046: 2, H0009: 2, H0081:	
			2. H0620: 2. H0266: 2.	
			H0124: 2, H0135: 2, H0551:	
			2 H0100: 2 1 0646: 2	
			1 0768. 2 1 0774. 2 1 0806.	
			LOVOS. 2, LOV /4: 2, LOSOS.	
			2, H0435: 2, H0539: 2,	
-			L0748: 2, L0754: 2, L0588:	
			2, L0589: 2, L0608: 2,	
			L0593: 2, H0543: 2, S0384:	
			2. H0170: 1. H0140: 1.	
			H0716: 1. H0650: 1. H0656:	
	,		1. H0254: 1. H0300: 1.	
	 		H0638: 1, S0410: 1, H0637:	
		·	1, S0045: 1, S0046: 1,	
			S0476: 1, H0619: 1, S0278:	
			1, S0222: 1, H0600: 1,	
			H0497: 1, H0632: 1, H0559:	
			1, H0013: 1, H0069: 1,	
			H0042: 1, H0706: 1, S0010:	
			1, S0182: 1, H0318: 1,	
			H0263: 1, T0110: 1, L0471:	
			1, H0024: 1, H0416: 1,	
			H0290: 1, H0292: 1, H0286:	
	 		1, S0250: 1, H0622: 1,	
			L0194: 1, L0483: 1, T0006:	
			1, H0213: 1, H0644: 1,	
			L0142: 1, H0181: 1, H0606:	
			1, L0055: 1, H0090: 1,	
			H0038: 1, H0616: 1, T0067:	
			1, H0488: 1, H0412: 1,	

H0056: 1, T0041: 1, T0042: 1, L0475: 1, H0396: 1, S0144: 1, S0142: 1, S0210: 1, S0002: 1, H0695: 1, L0763: 1, L0763: 1, L0770: 1, L0769: 1, L0662: 1, L0649: 1, L0381: 1, L0388: 1, L0573: 1, L0570: 1, L0573: 1, L0570: 1, L0571: 1, L0571: 1, L0571: 1, L0571: 1, L0571: 1, L0566: 1, L0517: 1, L0566: 1, H0698: 1, H0689: 1, H0684: 1, H0672: 1, S0330: 1, L0602: 1, H0672: 1, S0330: 1, L0602: 1, H0672: 1, S0330: 1, L0573: 1, L0778: 1, H0678: 1, S00078: 1, S00078: 1, L0599: 1, S0006: 1, H0673: 1, H0672: 1, S00078: 1, H0672: 1, S00078: 1, H0672: 1, H0673: 1, H0672: 1, S00078: 1, H0673: 1, H0672: 1, S00078: 1, H0672: 1, S00078: 1, H0672: 1, S00078: 1, H0673: 1, H0672: 1, S00078: 1, H0673: 1, H0672: 1, S00078: 1, H0673: 1, H0672: 1, S00078: 1, H0672: 1,	AR060: 2 H0600: 1	H0437: 2, H0587: 2, H0494: 2, L0769: 2, H0547: 2, S0028: 2, L0439: 2, L0593: 2, H0556: 1, H0657: 1, H0662: 1, H0125: 1, S0418: 1, H0619: 1, H0618: 1, H0253: 1, H0318: 1, H0052: 1, H0135: 1, H0529: 1, L0438: 1, H0539:
	Pro-17 to Ser-24.	Gln-25 to Leu-30.
	1011	1012
	288 - 362	200 - 400
	207	208
	794016	740778
	HWDAJ01	HWHPB78
	497	498

1, H0521: 1, S0037: 1, S0424: 1, H0506: 1 and H0008: 1	AR089: 10, AR060: 6 L0665: 10, L0754: 6, L0438: 5, L0751: 5, L0777: 5, L0758: 4, S0046: 3, H0213: 3, L0769: 3, L0667: 3, L0771: 3, L0662: 3, L0659: 3, H0539: 3, L0747: 3, L0777: 3, S0276: 3, S0418: 2, H0208: 2, S0045: 2, H0428: 2, H0424: 2, L0764: 2, L0768: 2, L0769: 2, L0764: 2, L0768: 2, L0759: 2, L0764: 2, L0768: 2, L0759: 2, L0768: 2, L0769: 1, L0767: 3, S0276: 3, S0404: 2, L0768: 2, L0768: 2, L0759: 2, L0768: 2, L0759: 2, L0769: 1, H0551: 1, H0579: 1, H0579: 1, H0580: 1, S0132: 1, H0561: 1, H0578: 1, H0070: 1, H0586: 1, H0408: 1, S0049: 1, L0471: 1, H0408: 1, L0163: 1, R0208: 1, H0079: 1, H0256: 1, H0079: 1, H0256: 1, H0079: 1, H0239: 1, H0569: 1, H0079: 1, H0271: 1, H0509: 1, H0641: 1, S0210: 1, H0059: 1, L0630: 1, L0637: 1, H0059: 1, L0630: 1, L0630
	Pro-3 to Ala-8.
	1013
	1015 - 1203
	609 609
	789854
	HYABC84
	9.4 9.5

1, L0761: 1, L0646: 1, L0643: 1, L0773: 1, L0650: 1, L0657: 1, L0793: 1, L0383: 1, L0790: 1, L0792: 1, L0664: 1, S0052: 1, H0691: 1, H0593: 1, H0435: 1, H0672: 1, H0696: 1, H0576: 1, L0748: 1, L0745: 1, L0750: 1, L0731: 1, H0707: 1, L0596: 1, L0591: 1, L0592: 1, L0593: 1, L0595: 1, H0667: 1, H0422: 1 and L0600: 1.	AR089: 10, AR060: 6 L0655: 10, L0754: 6, L0438: 5, L0751: 5, L0777: 5, L0752: 5, L0755: 4, L0758: 4, S0046: 3, H0213: 3, L0769: 3, L0667: 3, L0771: 3, L0662: 3, L0659: 3, H0539: 3, L0747: 3, L0757: 3, S0276: 3, S0418: 2, H0208: 2, S0045: 2, H0428: 2, H0424: 2, H0553: 2, H0412: 2, L0649: 2, L0669: 2, L0666: 2, L0668: 2, L0649: 2, L0764: 2, L0768: 2, L0769: 2, L0743: 2, L0744: 2, L0439: 2, L0756: 2, L0759: 2, L0485: 2, L0599: 2, S0040: 1, S0342: 1, T0049: 1, H0583: 1, H0657: 1, R0212: 1, H0580: 1, S0132: 1, H0261: 1, H0550: 1, H0370: 1, H0586: 1, H0333: 1, H0013: 1, H0575: 1, H0618:
	Pro-3 to Ala-8.
	1014
	1080 - 1268
	510
	865064
	HYABC84
	200

				1, S0049: 1, H0052: 1,		
	-			H0009: 1, L0471: 1, H0620:		
				1, L0163: 1, S0388: 1,	-	
				S0051: 1, T0010: 1, H0408:		
<u> </u>				1, H0239: 1, H0266: 1,		
		-		H0179: 1, H0271: 1, H0124:		
				1, S0366: 1, H0135: 1,		
				H0059: 1, T0042: 1, H0509:		
			_	1, H0641: 1, S0210: 1,		
				H0529: 1, L0639: 1, L0637:		
				1, L0761: 1, L0646: 1,		
				L0643: 1, L0773: 1, L0650:		
				1, L0657: 1, L0635: 1,		
_		-		L0383: 1, L0790: 1, L0792:		
				1, L0664: 1, S0052: 1,		
		-		H0691: 1, H0593: 1, H0435:		
				1, H0672: 1, H0696: 1,		
				H0576: 1, L0748: 1, L0745:		
		-		1, L0750: 1, L0731: 1,		
		.,		H0707: 1, L0596: 1, L0591:		
				1, L0592: 1, L0593: 1,		
		_		L0595: 1, H0667: 1, H0422:		
				1 and L0600: 1.		

- [81] The first column in Table 1B provides the gene number in the application corresponding to the clone identifier. The second column in Table 1B provides a unique "Clone ID NO:Z" for a cDNA clone related to each contig sequence disclosed in Table 1B. This clone ID references the cDNA clone which contains at least the 5' most sequence of the assembled contig and at least a portion of SEQ ID NO:X was determined by directly sequencing the referenced clone. The reference clone may have more sequence than described in the sequence listing or the clone may have less. In the vast majority of cases, however, the clone is believed to encode a full-length polypeptide. In the case where a clone is not full-length, a full-length cDNA can be obtained by methods described elsewhere herein.
- [82] The third column in Table 1B provides a unique "Contig ID" identification for each contig sequence. The fourth column provides the "SEQ ID NO:" identifier for each of the contig polynucleotide sequences disclosed in Table 1B. The fifth column, "ORF (From-To)", provides the location (i.e., nucleotide position numbers) within the polynucleotide sequence "SEQ ID NO:X" that delineate the preferred open reading frame (ORF) shown in the sequence listing and referenced in Table 1B, column 6, as SEQ ID NO:Y. Where the nucleotide position number "To" is lower than the nucleotide position number "From", the preferred ORF is the reverse complement of the referenced polynucleotide sequence.
- [83] The sixth column in Table 1B provides the corresponding SEQ ID NO:Y for the polypeptide sequence encoded by the preferred ORF delineated in column 5. In one embodiment, the invention provides an amino acid sequence comprising, or alternatively consisting of, a polypeptide encoded by the portion of SEQ ID NO:X delineated by "ORF (From-To)". Also provided are polynucleotides encoding such amino acid sequences and the complementary strand thereto.
- [84] Column 7 in Table 1B lists residues comprising epitopes contained in the polypeptides encoded by the preferred ORF (SEQ ID NO:Y), as predicted using the algorithm of Jameson and Wolf, (1988) Comp. Appl. Biosci. 4:181-186. The Jameson-

Wolf antigenic analysis was performed using the computer program PROTEAN (Version 3.11 for the Power MacIntosh, DNASTAR, Inc., 1228 South Park Street Madison, WI). In specific embodiments, polypeptides of the invention comprise, or alternatively consist of, at least one, two, three, four, five or more of the predicted epitopes as described in Table 1B. It will be appreciated that depending on the analytical criteria used to predict antigenic determinants, the exact address of the determinant may vary slightly.

[85] Column 8, in Table 1B, provides an expression profile and library code: count for each of the contig sequences (SEQ ID NO:X) disclosed in Table 1B, which can routinely be combined with the information provided in Table 4 and used to determine the tissues, cells, and/or cell line libraries which predominantly express the polynucleotides of the invention. The first number in column 8 (preceding the colon), represents the tissue/cell source identifier code corresponding to the code and description provided in Table 4. For those identifier codes in which the first two letters are not "AR", the second number in column 8 (following the colon) represents the number of times a sequence corresponding to the reference polynucleotide sequence was identified in the tissue/cell source. Those tissue/cell source identifier codes in which the first two letters are "AR" designate information generated using DNA array technology. Utilizing this technology, cDNAs were amplified by PCR and then transferred, in duplicate, onto the array. Gene expression was assayed through hybridization of first strand cDNA probes to the DNA array, cDNA probes were generated from total RNA extracted from a variety of different tissues and cell lines. Probe synthesis was performed in the presence of <sup>33</sup>P dCTP, using oligo(dT) to prime reverse transcription. After hybridization, high stringency washing conditions were employed to remove non-specific hybrids from the array. The remaining signal, emanating from each gene target, was measured using a Phosphorimager. Gene expression was reported as Phosphor Stimulating Luminescence (PSL) which reflects the level of phosphor signal generated from the probe hybridized to each of the gene targets represented on the array. A local background signal subtraction was performed before the total signal generated from each array was used to normalize gene expression between the different hybridizations. The value presented after "[array code]:" represents the mean of the duplicate values, following background subtraction and probe normalization. One of skill in the art could routinely use this information to identify normal and/or diseased tissue(s) which show a predominant expression pattern of the corresponding

polynucleotide of the invention or to identify polynucleotides which show predominant and/or specific tissue and/or cell expression.

[86] Column 9 in Table 1B provides a chromosomal map location for certain polynucleotides of the invention. Chromosomal location was determined by finding exact matches to EST and cDNA sequences contained in the NCBI (National Center for Biotechnology Information) UniGene database. Each sequence in the UniGene database is assigned to a "cluster"; all of the ESTs, cDNAs, and STSs in a cluster are believed to be derived from a single gene. Chromosomal mapping data is often available for one or more sequence(s) in a UniGene cluster; this data (if consistent) is then applied to the cluster as a whole. Thus, it is possible to infer the chromosomal location of a new polynucleotide sequence by determining its identity with a mapped UniGene cluster.

[87] A modified version of the computer program BLASTN (Altshul, et al., J. Mol. Biol. 215:403-410 (1990), and Gish, and States, Nat. Genet. 3:266-272) (1993) was used to search the UniGene database for EST or cDNA sequences that contain exact or near-exact matches to a polynucleotide sequence of the invention (the 'Query'). A sequence from the UniGene database (the 'Subject') was said to be an exact match if it contained a segment of 50 nucleotides in length such that 48 of those nucleotides were in the same order as found in the Query sequence. If all of the matches that met this criteria were in the same UniGene cluster, and mapping data was available for this cluster, it is indicated in Table 1B under the heading "Cytologic Band". Where a cluster had been further localized to a distinct cytologic band, that band is disclosed; where no banding information was available, but the gene had been localized to a single chromosome, the chromosome is disclosed.

[88] Once a presumptive chromosomal location was determined for a polynucleotide of the invention, an associated disease locus was identified by comparison with a database of diseases which have been experimentally associated with genetic loci. The database used was the Morbid Map, derived from OMIM<sup>TM</sup> ("Online Mendelian Inheritance in Man"; McKusick-Nathans Institute for Genetic Medicine, Johns Hopkins University (Baltimore, MD) and National Center for Biotechnology Information, National Library of Medicine (Bethesda, MD) 2000; World Wide Web URL: http://www.ncbi.nlm.nih.gov/omim/). If the putative chromosomal location of a polynucleotide of the invention (Query sequence)

was associated with a disease in the Morbid Map database, an OMIM reference identification number was noted in column 10, Table 1B, labelled "OMIM Disease Reference(s). Table 5 is a key to the OMIM reference identification numbers (column 1), and provides a description of the associated disease in Column 2.

[89] Table 1C summarizes additional polynucleotides encompassed by the invention (including cDNA clones related to the sequences (Clone ID NO:Z), contig sequences (contig identifier (Contig ID:) contig nucleotide sequence identifiers (SEQ ID NO:X)), and genomic sequences (SEQ ID NO:B). The first column provides a unique clone identifier, "Clone ID NO:Z", for a cDNA clone related to each contig sequence. The second column provides the sequence identifier, "SEO ID NO:X", for each contig sequence. The third column provides a unique contig identifier, "Contig ID:" for each contig sequence. The fourth column, provides a BAC identifier "BAC ID NO:A" for the BAC clone referenced in the corresponding row of the table. The fifth column provides the nucleotide sequence identifier, "SEQ ID NO:B" for a fragment of the BAC clone identified in column four of the corresponding row of the table. The sixth column, "Exon From-To", provides the location (i.e., nucleotide position numbers) within the polynucleotide sequence of SEO ID NO:B which delineate certain polynucleotides of the invention that are also exemplary members of polynucleotide sequences that encode polypeptides of the invention (e.g., polypeptides containing amino acid sequences encoded by the polynucleotide sequences delineated in column six, and fragments and variants thereof).

## TABLE 1C

Clone ID	SEQ ID No:X	CONTIG ID	BAC ID: A	SEQ ID NO:B	EXON From-To
H6BSF56	11	762968	AC069362	1019	1-131
H6BSF56	11	762968	AC027584	1020	1-162
H6BSF56	11	762968	AC011101	1021	1-100
H6BSF56	11	762968	AC073446	1022	1-140
H6BSF56	11	762968	AC026556	1023	1-114
H6BSF56	11	762968	AL136171	1024	1-61
H6BSF56	11	762968	AC025975	1025	1-136
H6BSF56	11	762968	AC073219	1026	1-123
H6BSF56	11	762968	AL162741	1027	1-45
H6BSF56	11	762968	AC027584	1028	1-368
H6BSF56	11	762968	AC073446	1029	1-52
					2626-2925
H6BSF56	11	762968	AL162741	1030	1-102
H6EEC72	13	889401	AC012314	1031	1-181
					1281-1463
			1		2719-2983
					3158-3411
					3804-6347
					6745-6879
					7118-7319
					7420-7521
					7859-8305
					8552-8602
					9988-10334
					10415-10778
					11003-11127
					11210-11303
					11334-11832
					13093-13145
				•	13703-13837
				·	13918-14152 15415-15511
		-			15613-15742
					15998-16087
				İ	16231-16307
					16447-17211
					18520-18796
					21777-22001
H6EEC72	13	889401	AC009968	1032	1-180
Hollec/2		005401	710005500	1032	1275-1457
					2712-2976
					3150-3403
					3796-6332
					6730-6864
					7103-7303
					7404-7505
					7843-8289
					8536-8586
					9970-10312
					10393-10756
					10981-11105
					11188-11805
	1			1	13068-13120

					13678-13812	
					13905-13994	
H6EEC72	13	889401	AC012314	1033	1-43	
	İ				861-1031	
						1576-1743
						1924-2132
				2473-2905		
				İ	3177-3360	
					3651-4332	
					4422-4583	
					4830-4995	
					5086-5365	
H6EEC72	13	889401	AC009968	1034	1-43	
Hobbers	10				857-1027	
					1570-1737	
		•			1918-2126	
					2197-2426	
					2467-2899	
					3171-3354	
					3644-4326	
					4416-4577	
]					4824-4989	
	İ				5080-5360	
HACAB68	14	584773	AL160283	1035	1-2811	
HACAB68	14	584773	AL354793	1036	1-3734	
HACAB06	14	304773	1112334733	1000	3843-4723	
TIACADO	14	584773	AL356058	1037	1-3055	
HACAB68	14	304773	ALSSOOSO	1037	3165-4045	
IIA CDISC	15	847112	AC069497	1038	1-117	
HACBJ56	13	04/112	ACOUPTI	1050	2470-3367	
						4908-5262
1					5641-5756	
					ļ	7886-8200
			:		9815-11138	
TIA CD ISC	1.5	847112	AC007104	1039	1-802	
HACBJ56	15	84/112	AC00/104	1039	2342-2695	
		1			3074-3189	
					5319-5633	
		,			7248-8571	
77. CD 75.	1.5	047112	AC069497	1040	1-453	
HACBJ56	15	847112	AC007104	1041	1-453	
HACBJ56	15	847112	AC007104 AC012073	1041	1-134	
HACBS22	16	847113	AC012073	1042	718-833	
	<u>,</u>				1002-1132	
					2357-2516	
					3762-3945	
					5344-5477	
					7446-7594	
					7742-7904	
					10636-10725	
					11138-12223	
					12583-12977	
					13095-13178	
					14224-14532	
1					14224-14332	
					15779-16124	
					16257-16343	
					16508-16826	
1					17489-17757	

					17847-18008
					19028-19192
					19755-23561
					24286-24717
					24920-25347
					25567-25741
					26629-26891
					27895-27968
HACBS22	16	847113	AC012073	1043	1-545
HADMB15	19	847116	AC026666	1044	1-385
HADNIDIS	13	047110	110020000	10	406-780
HADMB15	19	847116	AC026281	1045	1-114
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					20028-20295
HYABC84	510	865064	AL132825	2161	1-188

- [90] Tables 1D and 1E: The polynucleotides or polypeptides, or agonists or antagonists of the present invention can be used in assays to test for one or more biological activities. If these polynucleotides and polypeptides do exhibit activity in a particular assay, it is likely that these molecules may be involved in the diseases associated with the biological activity. Thus, the polynucleotides or polypeptides, or agonists or antagonists could be used to treat the associated disease.
- [91] The present invention encompasses methods of preventing, treating, diagnosing, or ameliorating a disease or disorder. In preferred embodiments, the present invention encompasses a method of treating a disease or disorder listed in the "Preferred Indications" columns of Table 1D and Table 1E; comprising administering to a patient in which such treatment, prevention, or amelioration is desired a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) in an amount effective to treat, prevent, diagnose, or ameliorate the disease or disorder. The first and second columns of Table 1D show the "Gene No." and "cDNA Clone ID No.", respectively, indicating certain nucleic acids and proteins (or antibodies against the same) of the invention (including polynucleotide, polypeptide, and antibody fragments or variants thereof) that may be used in preventing, treating, diagnosing, or ameliorating the disease(s) or disorder(s) indicated in the corresponding row in Column 3 of Table 1D.
- [92] In another embodiment, the present invention also encompasses methods of preventing, treating, diagnosing, or ameliorating a disease or disorder listed in the "Preferred Indications" column of Table 1D and Table 1E; comprising administering to a patient combinations of the proteins, nucleic acids, or antibodies of the invention (or fragments or variants thereof), sharing similar indications as shown in the corresponding rows in Column 3 of Table 1D.
- [93] The "Preferred Indications" columns of Table 1D and Table 1E describe diseases, disorders, and/or conditions that may be treated, prevented, diagnosed, or ameliorated by a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof).
- [94] The recitation of "Cancer" in the "Preferred Indications" columns indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof) may be used for example, to diagnose, treat, prevent, and/or

ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., leukemias, cancers, and/or as described below under "Hyperproliferative Disorders").

[95] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Cancer" recitation in the "Preferred Indication" column of Table 1D may be used for example, to diagnose, treat, prevent, and/or ameliorate a neoplasm located in a tissue selected from the group consisting of: colon, abdomen, bone, breast, digestive system, liver, pancreas, prostate, peritoneum, lung, blood (e.g., leukemia), endocrine glands (adrenal, parathyroid, pituitary, testicles, ovary, thymus, thyroid), uterus, eye, head and neck, nervous (central and peripheral), lymphatic system, pelvic, skin, soft tissue, spleen, thoracic, and urogenital.

[96] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Cancer" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a pre-neoplastic condition, selected from the group consisting of: hyperplasia (e.g., endometrial hyperplasia and/or as described in the section entitled "Hyperproliferative Disorders"), metaplasia (e.g., connective tissue metaplasia, atypical metaplasia, and/or as described in the section entitled "Hyperproliferative Disorders"), and/or dysplasia (e.g., cervical dysplasia, and bronchopulmonary dysplasia).

[97] In another specific embodiment, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Cancer" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a benign dysproliferative disorder selected from the group consisting of: benign tumors, fibrocystic conditions, tissue hypertrophy, and/or as described in the section entitled "Hyperproliferative Disorders".

[98] The recitation of "Immune/Hematopoietic" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), blood disorders (e.g., as described below under "Immune Activity" "Cardiovascular Disorders" and/or "Blood-Related Disorders"), and infections (e.g., as described below under "Infectious Disease").

[99] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having the "Immune/Hematopoietic" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat,

prevent, and/or ameliorate a disease or disorder selected from the group consisting of: anemia, pancytopenia, leukopenia, thrombocytopenia, leukemias, Hodgkin's disease, non-Hodgkin's lymphoma, acute lymphocytic anemia (ALL), plasmacytomas, multiple myeloma, Burkitt's lymphoma, arthritis, asthma, AIDS, autoimmune disease, rheumatoid arthritis, granulomatous disease, immune deficiency, inflammatory bowel disease, sepsis, neutropenia, neutrophilia, psoriasis, immune reactions to transplanted organs and tissues, systemic lupus erythematosis, hemophilia, hypercoagulation, diabetes mellitus, endocarditis, meningitis, Lyme Disease, and allergies.

[100] The recitation of "Reproductive" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), and disorders of the reproductive system (e.g., as described below under "Reproductive System Disorders").

[101] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Reproductive" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: cryptorchism, prostatitis, inguinal hernia, varicocele, leydig cell tumors, verrucous carcinoma, prostatitis, malacoplakia, Peyronie's disease, penile carcinoma, squamous cell hyperplasia, dysmenorrhea, ovarian adenocarcinoma, Turner's syndrome, mucopurulent cervicitis, Sertoli-leydig tumors, ovarian cancer, uterine cancer, pelvic inflammatory disease, testicular cancer, prostate cancer, Klinefelter's syndrome, Young's syndrome, premature ejaculation, diabetes mellitus, cystic fibrosis, Kartagener's syndrome, testicular atrophy, testicular feminization, anorchia, ectopic testis, epididymitis, orchitis, gonorrhea, syphilis, testicular torsion, vasitis nodosa, germ cell tumors, stromal tumors, dysmenorrhea, retroverted uterus, endometriosis, fibroids, adenomyosis, anovulatory bleeding, amenorrhea, Cushing's syndrome, hydatidiform moles, Asherman's syndrome, premature menopause, precocious puberty, uterine polyps, dysfunctional uterine bleeding, cervicitis, chronic cervicitis, mucopurulent cervicitis, cervical dysplasia, cervical polyps, Nabothian cysts, cervical erosion, cervical incompetence, cervical neoplasms, pseudohermaphroditism, and premenstrual syndrome.

[102] The recitation of "Musculoskeletal" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the

invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), and disorders of the immune system (e.g., as described below under "Immune Activity").

[103] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Musculoskeletal" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: bone cancers (e.g., osteochondromas, benign chondromas, chondroblastoma, chondromyxoid fibromas, osteoid osteomas, giant cell tumors, multiple myeloma, osteosarcomas), Paget's Disease, rheumatoid arthritis, systemic lupus erythematosus, osteomyelitis, Lyme Disease, gout, bursitis, tendonitis, osteoporosis, osteoarthritis, muscular dystrophy, mitochondrial myopathy, cachexia, and multiple sclerosis.

[104] The recitation of "Cardiovascular" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), and disorders of the cardiovascular system (e.g., as described below under "Cardiovascular Disorders").

[105] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Cardiovascular" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: myxomas, fibromas, rhabdomyomas, cardiovascular abnormalities (e.g., congenital heart defects, cerebral arteriovenous malformations, septal defects), heart disease (e.g., heart failure, congestive heart disease, arrhythmia, tachycardia, fibrillation, pericardial Disease, endocarditis), cardiac arrest, heart valve disease (e.g., stenosis, regurgitation, prolapse), vascular disease (e.g., hypertension, coronary artery disease, angina, aneurysm, arteriosclerosis, peripheral vascular disease), hyponatremia, hypernatremia, hypokalemia, and hyperkalemia.

[106] The recitation of "Mixed Fetal" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent,

and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders").

[107] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Mixed Fetal" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: spina bifida, hydranencephaly, neurofibromatosis, fetal alcohol syndrome, diabetes mellitus, PKU, Down's syndrome, Patau syndrome, Edwards syndrome, Turner syndrome, Apert syndrome, Carpenter syndrome, Conradi syndrome, Crouzon syndrome, cutis laxa, Cornelia de Lange syndrome, Ellis-van Creveld syndrome, Holt-Oram syndrome, Kartagener syndrome, Meckel-Gruber syndrome, Noonan syndrome, Pallister-Hall syndrome, Rubinstein-Taybi syndrome, Scimitar syndrome, Smith-Lemli-Opitz syndrome, thromocytopenia-absent radius (TAR) syndrome, Treacher Collins syndrome, Williams syndrome, Hirschsprung's disease, Meckel's diverticulum, polycystic kidney disease, Turner's syndrome, and gonadal dysgenesis, Klippel-Feil syndrome, Ostogenesis imperfecta, muscular dystrophy, Tay-Sachs disease, Wilm's tumor, neuroblastoma, and retinoblastoma.

[108] The recitation of "Excretory" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and renal disorders (e.g., as described below under "Renal Disorders").

[109] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Excretory" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: bladder cancer, prostate cancer, benign prostatic hyperplasia, bladder disorders (e.g., urinary incontinence, urinary retention, urinary obstruction, urinary tract Infections, interstitial cystitis, prostatitis, neurogenic bladder, hematuria), renal disorders (e.g., hydronephrosis, proteinuria, renal failure, pyelonephritis, urolithiasis, reflux nephropathy, and unilateral obstructive uropathy).

[110] The recitation of "Neural/Sensory" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the

invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and diseases or disorders of the nervous system (e.g., as described below under "Neural Activity and Neurological Diseases").

[111] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Neural/Sensory" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: brain cancer (e.g., brain stem glioma, brain tumors, central nervous system (Primary) lymphoma, central nervous system lymphoma, cerebellar astrocytoma, and cerebral astrocytoma, neurodegenerative disorders (e.g., Alzheimer's Disease, Creutzfeldt-Jakob Disease, Parkinson's Disease, and Idiopathic Presenile Dementia), encephalomyelitis, cerebral malaria, meningitis, metabolic brain diseases (e.g., phenylketonuria and pyruvate carboxylase deficiency), cerebellar ataxia, ataxia telangiectasia, and AIDS Dementia Complex, schizophrenia, attention deficit disorder, hyperactive attention deficit disorder, autism, and obsessive compulsive disorders.

[112] The recitation of "Respiratory" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and diseases or disorders of the respiratory system (e.g., as described below under "Respiratory Disorders").

[113] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Respiratory" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: cancers of the respiratory system such as larynx cancer, pharynx cancer, trachea cancer, epiglottis cancer, lung cancer, squamous cell carcinomas, small cell (oat cell) carcinomas, large cell carcinomas, and adenocarcinomas. Allergic reactions, cystic fibrosis, sarcoidosis, histiocytosis X, infiltrative lung diseases (e.g., pulmonary fibrosis and lymphoid interstitial pneumonia), obstructive airway diseases (e.g., asthma, emphysema, chronic or acute bronchitis), occupational lung diseases (e.g., silicosis and asbestosis), pneumonia, and pleurisy.

[114] The recitation of "Endocrine" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and diseases or disorders of the respiratory system (e.g., as described below under "Respiratory Disorders"), renal disorders (e.g., as described below under "Renal Disorders"), and disorders of the endocrine system (e.g., as described below under "Endocrine Disorders".

[115] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having an "Endocrine" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: cancers of endocrine tissues and organs (e.g., cancers of the hypothalamus, pituitary gland, thyroid gland, parathyroid glands, pancreas, adrenal glands, ovaries, and testes), diabetes (e.g., diabetes insipidus, type I and type II diabetes mellitus), obesity, disorders related to pituitary glands (e.g., hyperpituitarism, hypopituitarism, and pituitary dwarfism), hypothyroidism, hyperthyroidism, goiter, reproductive disorders (e.g. male and female infertility), disorders related to adrenal glands (e.g., Addison's Disease, corticosteroid deficiency, and Cushing's Syndrome), kidney cancer (e.g., hypernephroma, transitional cell cancer, and Wilm's tumor), diabetic nephropathy, interstitial nephritis, polycystic kidney disease, glomerulonephritis (e.g., IgM mesangial proliferative glomerulonephritis and glomerulonephritis caused by autoimmune disorders; such as Goodpasture's syndrome), and nephrocalcinosis.

[116] The recitation of "Digestive" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and diseases or disorders of the gastrointestinal system (e.g., as described below under "Gastrointestinal Disorders".

[117] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Digestive" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: ulcerative colitis, appendicitis, Crohn's disease, hepatitis, hepatic encephalopathy, portal hypertension,

cholelithiasis, cancer of the digestive system (e.g., biliary tract cancer, stomach cancer, colon cancer, gastric cancer, pancreatic cancer, cancer of the bile duct, tumors of the colon (e.g., polyps or cancers), and cirrhosis), pancreatitis, ulcerative disease, pyloric stenosis, gastroenteritis, gastritis, gastric atropy, benign tumors of the duodenum, distension, irritable bowel syndrome, malabsorption, congenital disorders of the small intestine, bacterial and parasitic infection, megacolon, Hirschsprung's disease, aganglionic megacolon, acquired megacolon, colitis, anorectal disorders (e.g., anal fistulas, hemorrhoids), congenital disorders of the liver (e.g., Wilson's disease, hemochromatosis, cystic fibrosis, biliary atresia, and alpha1-antitrypsin deficiency), portal hypertension, cholelithiasis, and jaundice.

[118] The recitation of "Connective/Epithelial" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), cellular and genetic abnormalities (e.g., as described below under "Diseases at the Cellular Level "), angiogenesis (e.g., as described below under "Anti-Angiogenesis Activity "), and or to promote or inhibit regeneration (e.g., as described below under "Regeneration "), and wound healing (e.g., as described below under "Wound Healing and Epithelial Cell Proliferation").

[119] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Connective/Epithelial" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: connective tissue metaplasia, mixed connective tissue disease, focal epithelial hyperplasia, epithelial metaplasia, mucoepithelial dysplasia, graft v. host disease, polymyositis, cystic hyperplasia, cerebral dysplasia, tissue hypertrophy, Alzheimer's disease, lymphoproliferative disorder, Waldenstron's macroglobulinemia, Crohn's disease, pernicious anemia, idiopathic Addison's disease, glomerulonephritis, bullous pemphigoid, Sjogren's syndrome, diabetes mellitus, cystic fibrosis, osteoblastoma, osteoclastoma, osteosarcoma, chondrosarcoma, osteoporosis, osteocarthritis, periodontal disease, wound polychondritis, vasculitis, polyarteritis Wegener's healing, relapsing nodosa, granulomatosis, cellulitis, rheumatoid arthritis, psoriatic arthritis, discoid lupus erythematosus, systemic lupus erythematosus, scleroderma, CREST syndrome, Sjogren's syndrome, polymyositis, dermatomyositis, mixed connective tissue disease, relapsing polychondritis, vasculitis, Henoch-Schonlein syndrome, erythema nodosum, polyarteritis nodosa, temporal (giant cell) arteritis, Takayasu's arteritis, Wegener's granulomatosis, Reiter's syndrome, Behcet's syndrome, ankylosing spondylitis, cellulitis, keloids, Ehler Danlos syndrome, Marfan syndrome, pseudoxantoma elasticum, osteogenese imperfecta, chondrodysplasias, epidermolysis bullosa, Alport syndrome, and cutis laxa.

**TABLE 1D** 

Gene No.	Clone ID	Preferred Indications
1	H6BSF56	Cancer
2	H6EDM64	Cancer
3	H6EEC72	Cancer
4	HACAB68	Connective/Epithelial,
		Immune/Hematopoetic
5	HACBJ56	Cancer
6	HACBS22	Cancer
7	HADDE71	Cancer
8	HADDJ13	Connective/Epithelial
9	HADMB15	Cancer
10	HAGBQ12	Excretory,
	`	Neural/Sensory
11	HAGDW20	Neural/Sensory,
		Reproductive
12	HAGEG10	Cancer
13	HAGEQ79	Cancer
14	HAGFS57	Cancer
15	HAGHN57	Cancer
16	HAHEA15	Cardiovascular
17	HAJAA47	Immune/Hematopoetic
18	HAJAY92	Cancer
19	HAJBV67	Cancer
20	HAJCH70	Cancer
21	HAOAG15	Cancer
22	HAQAI92	Digestive,
	,	Mixed Fetal,
		Reproductive
23	HAQCE11	Reproductive
24	HATBI94	Cancer
25	HATCB45	Endocrine,
		Immune/Hematopoetic
26	HATCD80	Endocrine,
		Reproductive
27	HATCI03	Endocrine,
		Immune/Hematopoetic,
		Neural/Sensory
28	HATEH20	Cancer
29	HBAGD86	Cancer
30	HBCJL35	Cancer
31	HBDAB91	Digestive,
		Immune/Hematopoetic
32	HBDAB91	Digestive,

	···	
		Immune/Hematopoetic
33	HBGBC29	Cancer
34	HBGNC72	Cancer
35	HBHAA05	Neural/Sensory
36	HBHAA81	Cardiovascular,
		Neural/Sensory
37	HBIAA59	Cancer
38	HBIAC29	Cancer
39	HBICW51	Digestive,
		Immune/Hematopoetic,
		Neural/Sensory
40	HBJAB02	Cancer
41	HBJAC65	Cancer
42	HBJBM12	Immune/Hematopoetic
43	HBJCR46	Cancer
44	HBJDS79	Cancer
45	HBJDW56	Immune/Hematopoetic
46	HBJEL16	Cancer
47	HBJFK45	Immune/Hematopoetic
48	HBJIG20	Cancer
49	HBJKD16	Cancer
50	HBMBM96	Cancer
51	HBMBX01	Cancer
52	HBMTM11	Cancer
53	HBMTX26	Immune/Hematopoetic
54	HBMTY48	Immune/Hematopoetic,
54	IIDIVII I VO	Reproductive
55	HBMUH74	Cardiovascular,
	IIBMO1174	Immune/Hematopoetic,
		Reproductive
56	HBMWE61	Immune/Hematopoetic
57	HBNAX40	Cancer
58	HBNBJ76	Cancer
59	HBQAB79	Neural/Sensory
60	HBQAC57	Neural/Sensory
61	HBSAK32	Cancer
62	HBXCM66	Cardiovascular,
02	IIDACMOO	Neural/Sensory,
		Reproductive
63	HBXCX15	Immune/Hematopoetic,
63	IIBACATS	Neural/Sensory
64	HCDCY76	Cancer
65	HCDDL48	Musculoskeletal
66	HCE1G78	Cancer
67	HCE2H52	Immune/Hematopoetic,
] "	110021192	Neural/Sensory,
		Reproductive
68	HCE3B04	Cancer
69	HCE5F78	Immune/Hematopoetic,
"	IICESI 76	Neural/Sensory
70	HCEDR26	Digestive,
'	IICLDRZU	Immune/Hematopoetic,
		Neural/Sensory
71	HCEEE79	Neural/Sensory
72	HCEEQ25	Mixed Fetal,
'-	TICLEQ23	Neural/Sensory
73	HCEEU18	Cancer
L / 3	I IICEECTO	1

74	HCEFZ82	Cancer
75	HCEGX05	Cancer
76	HCFLN88	Cancer
77	HCFLT90	Cancer
78	HCHAB84	Cancer
79	HCMSX51	Cancer
80	HCNCO11	Digestive
81	HCNSD29	Cardiovascular,
		Digestive,
S		Immune/Hematopoetic
82	HCQBH72	Digestive,
		Excretory,
		Immune/Hematopoetic
83	HCQCC96	Cancer
84	HCQCJ56	Cancer
85	HCQCM24	Cancer
86	HCRAY10	Cancer
87	HCRBF72	Cancer
88	HCRNF78	Cancer
89	HCUAF85	Immune/Hematopoetic
90	HCUCF89	Immune/Hematopoetic
91	HCUCK44	Cancer
92	HCUDD64	Cancer
93	HCWAE64	Immune/Hematopoetic
94	HCWFU39	Endocrine,
		Immune/Hematopoetic,
		Neural/Sensory
95	HCWUL09	Immune/Hematopoetic,
	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	Neural/Sensory
96	HDHAA42	Cancer
97	HDHEB76	Cancer
98	HDPCW16	Cancer
99	HDPDI72	Digestive,
100	HDDD150	Immune/Hematopoetic
100	HDPDJ58	Cancer
101	HDPFF10	Cancer
102	HDPFV18	Cancer
103	HDPFY18	Cancer
104	HDPGE24	Cancer
105	HDPIU94	Cancer Cancer
106	HDPOC24 HDPOL37	Immune/Hematopoetic,
107	IDFOLS/	Reproductive
108	HDPOO76	Cancer
109	HDPPD93	Cancer
110	HDPPQ30	Immune/Hematopoetic
111	HDPPW82	Immune/Hematopoetic
1112	HDPXN20	Immune/Hematopoetic
113	HDQHM36	Immune/Hematopoetic
113	HDTAU35	Immune/Hematopoetic
115	HDTAV54	Cancer
	HDTFX18	Immune/Hematopoetic,
116		Reproductive
117	HDTGW48	Immune/Hematopoetic,
110	HDTI M10	Reproductive
118	HDTLM18 HE2CA60	Immune/Hematopoetic Cancer
	I HPZLANO	i Capret

120	THEOCACO	
120	HE2CA60	Cancer
121	HE2CH58	Digestive,
122	TIE2CN (20	Mixed Fetal
122	HE2CM39	Cancer
123	HE2HC60	Cancer
124	HE2PO93	Cancer
125	HE6AU52	Mixed Fetal
126	HE6CS65	Cancer
127	HE6DO92	Immune/Hematopoetic,
		Mixed Fetal
128	HE6EY13	Cancer
129	HE6FU11	Mixed Fetal,
		Neural/Sensory,
		Respiratory
130	HE6FV29	Cancer
131	HE8FC45	Cancer
132	HE8FC45	Cancer
133	HE8FD92	Cancer
134	HE8FD92	Cancer
135	HE8FD92	Cancer
136	HE8FD92	Cancer
137	HE8FD92	Cancer
138	HE8SG96	Mixed Fetal,
		Musculoskeletal
139	HE8TY46	Cancer
140	HE9CY05	Mixed Fetal
141	HE9EA10	Cancer
142	HE9GG20	Cancer
143	HEBCI18	Cancer
144	HEBCY54	Cancer
145	HEBDF77	Neural/Sensory
146	HEBDQ91	Neural/Sensory
147	HEBFR46	Cancer
148	HEBGE07	Neural/Sensory
149	HEGAU15	Excretory,
	112011011	Immune/Hematopoetic,
		Reproductive
150	HELAT35	Cardiovascular,
		Mixed Fetal
151	HELBU54	Cardiovascular
152	HELGG84	Cancer
153	HELGG84	Cancer
154	HEMEY47	Cardiovascular
155	HEOMC46	Immune/Hematopoetic
156	HEPBA14	Reproductive
157	HEQAH80	Cancer
158	HEQBF89	Reproductive
159	HETCI16	Cancer
160	HETDW58	Cancer
161	HETEY67	Connective/Epithelial,
101	HEIEIO/	Reproductive
162	HFCDW95	Cancer
163	HFCEI04	Neural/Sensory
164	HFCFD04	Cancer
165	HFCFE20	Cancer
166	HFEAY59	
167	HFGAJ16	Connective/Epithelial
10/	nrualio	Cancer

168	HFIHZ75	Cancer
169	HFIJA29	Cancer
170	HFIJA68	Cancer
171	HFKES05	Cancer
172	HFKEU12	Excretory
173	HFPCZ55	Cancer
174	HFPDR62	Immune/Hematopoetic,
		Neural/Sensory
175	HFPDS07	Cancer
176	HFRAB10	Excretory,
		Immune/Hematopoetic,
		Neural/Sensory
177	HFTBM38	Cancer
178	HFTDH56	Cancer
179	HFVGK35	Cancer
180	HFVHW43	Digestive
181	HFXAV37	Immune/Hematopoetic,
		Neural/Sensory
182	HFXBN86	Neural/Sensory
183	HFXBT66	Neural/Sensory
184	HFXFZ46	Neural/Sensory
185	HGBER72	Cancer
186	HGBEY14	Cancer
187	HGBGN34	Connective/Epithelial,
167	nobon34	
		Digestive, Reproductive
188	HGBHP91	Digestive
189	HGCAC19	
190		Cancer
	HGCAC19	Cancer
191	HGCAC19	Cancer
192	HHEAK45	Cancer
193	HHEGS55	Immune/Hematopoetic
194	HHEOW19	Cancer
195	HHFFF87	Cancer
196	HHFFL34	Cancer
197	HHFFS40	Cancer
198	HHGCS78	Immune/Hematopoetic
199	HHGDT26	Immune/Hematopoetic,
		Reproductive
200	HHPFU28	Cancer
201	HHPSA85	Cancer
202	HHSBI06	Cancer
203	HHSBI65	Cancer
204	HHSDI53	Cancer
205	HHSFC09	Cancer
206	HHSGL28	Cancer
207	HILCA24	Digestive,
		Immune/Hematopoetic,
		Reproductive
208	HILCA24	Digestive,
		Immune/Hematopoetic,
		Reproductive
209	HISAT67	Cancer
210	HJBCU75	Cancer
211	HJMAA03	Cancer
212	HJMAV41	Cancer
213	HJMAY90	Cancer
410	113IVIA I 90	Cancel

214	HJPBE39	Cancer
215	HJPBK28	Cancer
216	НЈРСН08	Cancer
217	HKABU43	Cancer
218	HKACI79	Cancer
219	HKAFF50	Cancer
220	HKGBF25	Cancer
221	HKIXC44	Cancer
222	HKMLK03	Digestive,
		Excretory,
		Immune/Hematopoetic
223	HKMLM95	Cancer
224	HKTAB41	Digestive,
		Excretory
225	HLDBG17	Cancer
226	HLDCA54	Cancer
227	HLDQU79	Cancer
228	HLDRT09	Cancer
229	HLHAP05	Immune/Hematopoetic,
		Neural/Sensory,
		Respiratory
230	HLHCS23	Respiratory
231	HLIBO72	Digestive
232	HLICE88	Digestive,
		Mixed Fetal
233	HLICO10	Cancer
234	HLJBS28	Cancer
235	HLMBW89	Cancer
236	HLMGP50	Digestive, Immune/Hematopoetic
237	HLMJB64	Cancer
238	HLMMX62	Immune/Hematopoetic,
		Neural/Sensory,
		Reproductive
239	HLQAS12	Cancer
240	HLQCL64	Cancer
241	HLQCX36	Digestive
242	HLWAF06	Digestive,
		Immune/Hematopoetic,
		Reproductive
243	HLWAU42	Cancer
244	HLWAU42	Cancer
245	HLWAV47	Cancer
246	HLWBB73	Cancer
247	HLWCN37	Cancer
248	HLWDB73	Cancer
249	HLYDF73	Immune/Hematopoetic
250	HLYEU59	Immune/Hematopoetic
251	HLYGB19	Cancer
252	HLYGE16	Cancer
253	HLYGY91	Cancer
254	HMCAZ04	Cancer
255	HMCAZ04	Cancer
256	HMCAZ04	Cancer
257	HMCAZ04	Cancer
258	HMCAZ04	Cancer
259	HMCFH60	Cancer

	T	1
260	HMDAB29	Digestive,
		Neural/Sensory
261	HMDAD44	Connective/Epithelial,
		Immune/Hematopoetic,
		Neural/Sensory
262	HMEBB82	Cancer
263	HMEDE24	Cancer
264	HMEDI90	Cancer
265	HMELM75	Cancer
266	HMIAK10	Neural/Sensory
267	HMIBF07	Neural/Sensory
268	HMICI80	Cardiovascular,
		Endocrine,
		Neural/Sensory
269	HMICP65	Cancer
270	HMJAK70	Neural/Sensory
271	HMSBE04	Immune/Hematopoetic
272	HMSCL38	Digestive,
		Immune/Hematopoetic,
,		Neural/Sensory
273	HMSCR69	Cancer
274	HMSHC86	Immune/Hematopoetic
275	HMSHU20	Immune/Hematopoetic,
		Reproductive
276	HMSHY25	Immune/Hematopoetic
277	HMTAB77	Cancer
278	HMUAE26	Cancer
279	HMUAN45	Cancer
280	HMVBC31	Cancer
281	HMVDU15	Cancer
282	HMWBL03	Cancer
283	HMWJF53	Cancer
284	HNEAK81	Immune/Hematopoetic
285	HNECL22	Cancer
286	HNECW49	Immune/Hematopoetic
287	HNEDH88	Immune/Hematopoetic
288	HNFAC50	Cancer
289	HNFGR08	Immune/Hematopoetic
290	HNFHF34	Cancer
291	HNGAK51	Immune/Hematopoetic
	HNGAM58	
292		Immune/Hematopoetic Immune/Hematopoetic
293	HNGBH53	<u> </u>
294	HNGDQ38	Immune/Hematopoetic
295	HNGDX18	Cancer
296	HNGDY34	Immune/Hematopoetic
297	HNGEA34	Digestive,
	1	Immune/Hematopoetic
298	HNGEQ75	Immune/Hematopoetic,
200	1000000	Neural/Sensory
299	HNGGA68	Immune/Hematopoetic,
	1	Musculoskeletal
300	HNGGP65	Immune/Hematopoetic
301	HNGHZ69	Immune/Hematopoetic
302	HNGIV64	Immune/Hematopoetic
303	HNGJB41	Immune/Hematopoetic
304	HNGKT41	Immune/Hematopoetic
305	HNGMW45	Immune/Hematopoetic

306	HNGNK44	Immune/Hematopoetic
307	HNGNO53	Immune/Hematopoetic
308	HNGPJ25	Immune/Hematopoetic,
		Mixed Fetal,
		Musculoskeletal
309	HNHEN82	Immune/Hematopoetic
310	HNHFE71	Immune/Hematopoetic
311	HNHGK22	Immune/Hematopoetic
312	HNHHB10	Immune/Hematopoetic,
		Reproductive
313	HNHKS19	Immune/Hematopoetic,
		Reproductive
314	HNTBT17	Cancer
315	HNTMH79	Cancer
316	HOABP31	Cancer
317	HOABP31	Cancer
318	HOACG07	Cancer
319	HODAG07	Reproductive
320	HODBB70	Reproductive
321	HODBV05	Cancer
322	HODCZ32	Reproductive
323	HOEBK60	Cancer
324	HOFAA78	Reproductive
325	HOFNB74	Reproductive
326	HOFNU55	Reproductive
327	HOGBF01	Reproductive
328	HORBS82	Cancer
329	HORBV76	Cardiovascular,
		Immune/Hematopoetic,
		Reproductive
330	HOSDO75	Cancer
331	HOSEC25	Immune/Hematopoetic,
		Musculoskeletal,
		Reproductive
332	HOSEI81	Digestive,
		Musculoskeletal
333	HOSEJ94	Cancer
334	HOUCA21	Connective/Epithelial,
		Immune/Hematopoetic,
		Musculoskeletal
335	HOUDE92	Cancer
336	HOUDR07	Cancer
337	HOUED72	Connective/Epithelial
338	HOUFS04	Cancer
339	HOUHI25	Cancer
340	HOVBD85	Musculoskeletal,
		Reproductive
341	HPCAB41	Immune/Hematopoetic,
		Reproductive
342	HPCAL26	Cancer
343	HPEAD23	Cancer
344	HPFBA54	Reproductive
345	HPFCI36	Cancer
346	HPFDI37	Cancer
347	HPIAA80	Cancer
348	HPJBJ51	Cancer
349	HPJBJ51	Cancer

351			
352	350	HPJBU43	Reproductive
353			
3554   HPMCV30   Cancer			
355			
356			Cancer
357		HPMFH77	Cancer
SSS	356	HPQAX38	Cardiovascular
359	357	HPQAX38	Cardiovascular
360	358	HPQCB83	Cancer
MPRCA64	359	HPQCC53	Cancer
362	360	HPRBH85	Cancer
363	361	HPRCA64	Cancer
364         HPWBA29         Reproductive           365         HPWDK06         Cancer           366         HRAAD30         Cancer           367         HRADA42         Cancer           368         HRADF49         Cancer           369         HRADN25         Cancer           370         HRADT25         Digestive, Excretory           371         HRDA117         Cancer           372         HRDDQ39         Cancer           373         HRDER22         Cancer           374         HRDEX93         Cancer           375         HRDFK37         Cancer           376         HRGBD54         Cancer           377         HROEA08         Cancer           378         HSAVA08         Immune/Hematopoetic           379         HSAVW42         Cancer           380         HSAWN53         Immune/Hematopoetic           381         HSAWZ40         Excretory, Immune/Hematopoetic, Reproductive           383         HSDEM54         Cancer           384         HSHBF76         Cancer           385         HSIFG47         Digestive           386         HSJBY32         Immune/He	362	HPRCD35	Cancer
According to the content of the co	363	HPTRM02	Cancer
365	364	HPWBA29	Reproductive
366         HRAAD30         Cancer           367         HRADA42         Cancer           368         HRADF49         Cancer           369         HRADN25         Cancer           370         HRADT25         Digestive,           370         HRADT25         Digestive,           370         HRADT25         Digestive,           370         HRDA117         Cancer           371         HRDA117         Cancer           372         HRDDQ39         Cancer           374         HRDEX22         Cancer           374         HRDEX93         Cancer           376         HRGBD54         Cancer           377         HROEA08         Cancer           378         HSAVA08         Immune/Hematopoetic           381         HSAWW53         Immune/Hematopoetic           381         HSAWX40         Immune/Hematopoetic,           382         HSAYC41         Excretory,           383         HSIFG47         Digestive           384         HSHBF76         Cancer           385         HSIFG47         Digestive           386         HSJBY32         Immune/Hematopoetic,	365	HPWDK06	
367	366		Cancer
368         HRADF49         Cancer           369         HRADN25         Cancer           370         HRADT25         Digestive, Excretory           371         HRDAI17         Cancer           372         HRDDQ39         Cancer           373         HRDER22         Cancer           374         HRDEX93         Cancer           375         HRDFK37         Cancer           376         HRGBD54         Cancer           377         HROEA08         Cancer           378         HSAVA08         Immune/Hematopoetic           379         HSAVW42         Cancer           380         HSAWN53         Immune/Hematopoetic           381         HSAYC41         Excretory, Immune/Hematopoetic, Reproductive           382         HSAYC41         Excretory, Immune/Hematopoetic, Musculoskeletal           384         HSHBF76         Cancer           385         HSIFG47         Digestive           386         HSJBY32         Immune/Hematopoetic, Musculoskeletal, Neural/Sensory           387         HSKDR27         Cancer           388         HSLH315         Musculoskeletal           390         HSNAP85         Cancer <td></td> <td></td> <td></td>			
369			
370			
Excretory			
371         HRDAI17         Cancer           372         HRDDQ39         Cancer           373         HRDER22         Cancer           374         HRDEX93         Cancer           375         HRDFK37         Cancer           376         HRGBD54         Cancer           377         HROEA08         Cancer           378         HSAVA08         Immune/Hematopoetic           379         HSAVW42         Cancer           380         HSAWN53         Immune/Hematopoetic           381         HSAWZ40         Immune/Hematopoetic           382         HSAYC41         Excretory, Immune/Hematopoetic, Reproductive           383         HSDZM54         Cancer           384         HSHBF76         Cancer           385         HSIFG47         Digestive           386         HSJBY32         Immune/Hematopoetic, Musculoskeletal, Neural/Sensory           387         HSKDR27         Cancer           388         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive	3,0	11101123	
372         HRDDQ39         Cancer           373         HRDER22         Cancer           374         HRDEX93         Cancer           375         HRDFK37         Cancer           376         HRGBD54         Cancer           377         HROEA08         Cancer           378         HSAVA08         Immune/Hematopoetic           379         HSAVW42         Cancer           380         HSAWN53         Immune/Hematopoetic           381         HSAWZ40         Immune/Hematopoetic           382         HSAYC41         Excretory, Immune/Hematopoetic, Reproductive           383         HSDZM54         Cancer           384         HSHBF76         Cancer           385         HSIFG47         Digestive           386         HSJBY32         Immune/Hematopoetic, Musculoskeletal, Neural/Sensory           387         HSKDR27         Cancer           388         HSLHG78         Cancer           389         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive	371	HRDAI17	
373         HRDER22         Cancer           374         HRDEX93         Cancer           375         HRDFK37         Cancer           376         HRGBD54         Cancer           377         HROEA08         Cancer           378         HSAVA08         Immune/Hematopoetic           379         HSAVW42         Cancer           380         HSAWN53         Immune/Hematopoetic           381         HSAWZ40         Immune/Hematopoetic           382         HSAYC41         Excretory, Immune/Hematopoetic, Reproductive           383         HSDZM54         Cancer           384         HSHBF76         Cancer           385         HSIFG47         Digestive           386         HSJBY32         Immune/Hematopoetic, Musculoskeletal, Neural/Sensory           387         HSKDR27         Cancer           388         HSLHG78         Cancer           389         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive			
374         HRDEX93         Cancer           375         HRDFK37         Cancer           376         HRGBD54         Cancer           377         HROEA08         Cancer           378         HSAVA08         Immune/Hematopoetic           379         HSAVW42         Cancer           380         HSAWN53         Immune/Hematopoetic           381         HSAWZ40         Immune/Hematopoetic           382         HSAYC41         Excretory, Immune/Hematopoetic, Reproductive           383         HSDZM54         Cancer           384         HSHBF76         Cancer           385         HSIFG47         Digestive           386         HSJBY32         Immune/Hematopoetic, Musculoskeletal, Neural/Sensory           387         HSKDR27         Cancer           388         HSLHG78         Cancer           389         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive           393         HSOAH16         Digestive			
375         HRDFK37         Cancer           376         HRGBD54         Cancer           377         HROEA08         Cancer           378         HSAVA08         Immune/Hematopoetic           379         HSAVW42         Cancer           380         HSAWN53         Immune/Hematopoetic           381         HSAWZ40         Immune/Hematopoetic           382         HSAYC41         Excretory, Immune/Hematopoetic, Reproductive           383         HSDZM54         Cancer           384         HSHBF76         Cancer           385         HSIFG47         Digestive           386         HSJBY32         Immune/Hematopoetic, Musculoskeletal, Neural/Sensory           387         HSKDR27         Cancer           388         HSLHG78         Cancer           389         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive           393         HSOAH16         Digestive			
376         HRGBD54         Cancer           377         HROEA08         Cancer           378         HSAVA08         Immune/Hematopoetic           379         HSAVW42         Cancer           380         HSAWN53         Immune/Hematopoetic           381         HSAWZ40         Immune/Hematopoetic           382         HSAYC41         Excretory, Immune/Hematopoetic, Reproductive           383         HSDZM54         Cancer           384         HSHBF76         Cancer           385         HSIFG47         Digestive           386         HSJBY32         Immune/Hematopoetic, Musculoskeletal, Neural/Sensory           387         HSKDR27         Cancer           388         HSLHG78         Cancer           389         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive           393         HSOAH16         Digestive			
377         HROEA08         Cancer           378         HSAVA08         Immune/Hematopoetic           379         HSAVW42         Cancer           380         HSAWN53         Immune/Hematopoetic           381         HSAWZ40         Immune/Hematopoetic           382         HSAYC41         Excretory, Immune/Hematopoetic, Reproductive           383         HSDZM54         Cancer           384         HSHBF76         Cancer           385         HSIFG47         Digestive           386         HSJBY32         Immune/Hematopoetic, Musculoskeletal, Neural/Sensory           387         HSKDR27         Cancer           388         HSLHG78         Cancer           389         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive           393         HSOAH16         Digestive			
378         HSAVA08         Immune/Hematopoetic           379         HSAVW42         Cancer           380         HSAWN53         Immune/Hematopoetic           381         HSAWZ40         Immune/Hematopoetic           382         HSAYC41         Excretory, Immune/Hematopoetic, Reproductive           383         HSDZM54         Cancer           384         HSHBF76         Cancer           385         HSIFG47         Digestive           386         HSJBY32         Immune/Hematopoetic, Musculoskeletal, Neural/Sensory           387         HSKDR27         Cancer           388         HSLHG78         Cancer           389         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive           393         HSOAH16         Digestive			
379         HSAVW42         Cancer           380         HSAWN53         Immune/Hematopoetic           381         HSAWZ40         Immune/Hematopoetic           382         HSAYC41         Excretory,           Immune/Hematopoetic,         Reproductive           383         HSDZM54         Cancer           384         HSHBF76         Cancer           385         HSIFG47         Digestive           386         HSJBY32         Immune/Hematopoetic,           Musculoskeletal,         Neural/Sensory           387         HSKDR27         Cancer           388         HSLHG78         Cancer           389         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive           393         HSOAH16         Digestive			
380         HSAWN53         Immune/Hematopoetic           381         HSAWZ40         Immune/Hematopoetic           382         HSAYC41         Excretory,           Immune/Hematopoetic,         Reproductive           383         HSDZM54         Cancer           384         HSHBF76         Cancer           385         HSIFG47         Digestive           386         HSJBY32         Immune/Hematopoetic,           Musculoskeletal,         Neural/Sensory           387         HSKDR27         Cancer           388         HSLHG78         Cancer           389         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive           393         HSOAH16         Digestive			
381         HSAWZ40         Immune/Hematopoetic           382         HSAYC41         Excretory, Immune/Hematopoetic, Reproductive           383         HSDZM54         Cancer           384         HSHBF76         Cancer           385         HSIFG47         Digestive           386         HSJBY32         Immune/Hematopoetic, Musculoskeletal, Neural/Sensory           387         HSKDR27         Cancer           388         HSLHG78         Cancer           389         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive           393         HSOAH16         Digestive			
382         HSAYC41         Excretory, Immune/Hematopoetic, Reproductive           383         HSDZM54         Cancer           384         HSHBF76         Cancer           385         HSIFG47         Digestive           386         HSJBY32         Immune/Hematopoetic, Musculoskeletal, Neural/Sensory           387         HSKDR27         Cancer           388         HSLHG78         Cancer           389         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive           393         HSOAH16         Digestive			<u> </u>
Immune/Hematopoetic,   Reproductive			Anna anna anna anna anna anna anna anna
Reproductive           383         HSDZM54         Cancer           384         HSHBF76         Cancer           385         HSIFG47         Digestive           386         HSJBY32         Immune/Hematopoetic, Musculoskeletal, Neural/Sensory           387         HSKDR27         Cancer           388         HSLHG78         Cancer           389         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive           393         HSOAH16         Digestive	302	113A 1 C41	
383         HSDZM54         Cancer           384         HSHBF76         Cancer           385         HSIFG47         Digestive           386         HSJBY32         Immune/Hematopoetic,			
384         HSHBF76         Cancer           385         HSIFG47         Digestive           386         HSJBY32         Immune/Hematopoetic, Musculoskeletal, Neural/Sensory           387         HSKDR27         Cancer           388         HSLHG78         Cancer           389         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive           393         HSOAH16         Digestive	282	HSD7M54	
385         HSIFG47         Digestive           386         HSJBY32         Immune/Hematopoetic, Musculoskeletal, Neural/Sensory           387         HSKDR27         Cancer           388         HSLHG78         Cancer           389         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive           393         HSOAH16         Digestive			
386         HSJBY32         Immune/Hematopoetic, Musculoskeletal, Neural/Sensory           387         HSKDR27         Cancer           388         HSLHG78         Cancer           389         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive           393         HSOAH16         Digestive		·····	
Musculoskeletal, Neural/Sensory           387         HSKDR27         Cancer           388         HSLHG78         Cancer           389         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive           393         HSOAH16         Digestive			
Neural/Sensory           387         HSKDR27         Cancer           388         HSLHG78         Cancer           389         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive           393         HSOAH16         Digestive	300	nsjb i sz	
387         HSKDR27         Cancer           388         HSLHG78         Cancer           389         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive           393         HSOAH16         Digestive			
388         HSLHG78         Cancer           389         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive           393         HSOAH16         Digestive	297	HCKDD27	
389         HSLHX15         Musculoskeletal           390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive           393         HSOAH16         Digestive			
390         HSNAP85         Cancer           391         HSNAZ09         Cancer           392         HSNBM34         Digestive           393         HSOAH16         Digestive			
391 HSNAZ09 Cancer 392 HSNBM34 Digestive 393 HSOAH16 Digestive			ii
392 HSNBM34 Digestive 393 HSOAH16 Digestive			
393 HSOAH16 Digestive			
			<u> </u>
394 HSQBF66 Cancer			
395 HSQDO85 Cancer			
396 HSQES57 Cancer			
397 HSRBE06 Cancer			
398 HSSDI26 Musculoskeletal			
399 HSSEA64 Cancer			Cancer
400 HSSEF77 Cancer			Cancer
401 HSSFE38 Cancer	401	HSSFE38	Cancer

402	HCCLEO	Margarianianiani
	HSSGJ58	Musculoskeletal
403	HSWBE76	Cancer
404	HSXCP38	Cardiovascular,
405	HSYBI06	Neural/Sensory Cancer
406	HT1SC27	
400	HT1SC27	Digestive,
		Immune/Hematopoetic, Reproductive
407	HT3BF49	Immune/Hematopoetic
408	HT4FV41	Cancer
409	HT5FX79	Cancer
410	HT5GR59	Cancer
411	HTAEI78	
412	HTDAA78	Immune/Hematopoetic
413		Cancer
413	HTEAG62	Digestive,
		Immune/Hematopoetic, Reproductive
414	HTECB02	Cancer
415	HTECEU2	Cancer
416	HTEDF18	Reproductive
417	HTEDJ28	Cancer
418	HTEDS12	Cardiovascular,
410	HIEDSIZ	Immune/Hematopoetic,
		Reproductive
419	HTEED26	Cancer
420	HTEED26	Cancer
421	HTEEF26	Cancer
422	HTEEF26	Cancer
423	HTEEW69	Reproductive
424	HTEGS07	Reproductive
425	HTEGS07	Cancer
426	HTEHA56	Cancer
427	HTEHU59	Cancer
428	HTEJD29	Reproductive
429	HTEKM46	Cancer
430	HTEMQ17	Cancer
431	HTENR63	Cancer
432	HTGGM44	Immune/Hematopoetic,
432	I I I GGM44	Musculoskeletal
433	HTHBZ06	Cancer
434	HTLAP64	Cancer
435	HTLBT80	Cancer
436	HTLDA84	Reproductive
437	HTLDN29	Cancer
438	HTLDU78	Reproductive
439	HTLEC82	Cancer
440	HTLEM16	Cancer
441	HTLEV48	Reproductive
442	HTLFA13	Musculoskeletal,
,72	IIILIAIS	Reproductive
443	HTLFI73	Cancer
444	HTLGI89	Cancer
445	HTLIF11	Cancer
446	HTLIF12	Excretory,
7-10	III LII 12	Reproductive
447	HTLIF12	Excretory,
		Reproductive
		1 F

		1 =
448	HTLIF12	Excretory,
		Reproductive
449	HTLIF12	Excretory,
		Reproductive
450	HTLIF12	Excretory,
		Reproductive
451	HTLIF12	Excretory,
		Reproductive
452	HTNAM63	Endocrine
453	HTNBK13	Cancer
454	HTOAI50	Immune/Hematopoetic
455	HTOAM11	Immune/Hematopoetic,
		Neural/Sensory
456	HTODH57	Immune/Hematopoetic
457	HTODH83	Immune/Hematopoetic
458	HTOEV16	Cancer
459	HTOGR38	Immune/Hematopoetic
460	НТОНО21	Immune/Hematopoetic
461	HTOHQ05	Immune/Hematopoetic
462	HTOJL95	Cancer
463	HTOJL95	Cancer
464	HTPDU17	Cancer
465	HTSFJ32	Immune/Hematopoetic
466	HTTCB60	Cancer
467	HTTEE41	Cancer
468	HTTEZ02	Cancer
469	HTWEH94	Immune/Hematopoetic
470	HTXBD09	Cancer
471	HTXDB22	Cancer
472	HTXDC38	Cancer
473	HTXDC38	Cancer
474	HTXDD61	
475		Cancer
476	HTXDG92 HTXET11	Cancer
		Immune/Hematopoetic
477	HTXFA72	Immune/Hematopoetic
478	HTXJY08	Cancer
479	HTXKF95	Cancer
480	HTXMZ07	Cancer
481	HUFCL31	Digestive,
		Immune/Hematopoetic
482	HUKBT67	Cancer
483	HUKDF20	Cardiovascular,
		Reproductive
484	HUKDY82	Cancer
485	HUSCJ14	Cancer
486	HUSGL67	Cancer
487	HUSGU40	Cancer
488	HUSIR18	Cancer
489	HUVDJ48	Digestive,
		Reproductive
490	HWAAI12	Cancer
491	HWBBQ70	Immune/Hematopoetic,
		Neural/Sensory
492	HWBCN36	Immune/Hematopoetic
493	HWBDJ08	Cancer
493 494	HWBDJ08 HWBFX16	Cancer Immune/Hematopoetic

		Immune/Hematopoetic, Neural/Sensory	
496	HWDAG96	Cancer	
497	HWDAJ01	Connective/Epithelial	
498	HWHPB78	Cancer	
499	HYABC84	Cancer	
500	HYABC84	Cancer	

[120] Table 1E provides information related to biological activities and preferred indications for polynucleotides and polypeptides of the invention (including antibodies, agonists, and/or antagonists thereof). Table 1E also provides information related to assays which may be used to test polynucleotides and polypeptides of the invention (including antibodies, agonists, and/or antagonists thereof) for the corresponding biological activities. The first column ("Gene No.") provides the gene number in the application for each clone identifier. The second column ("cDNA Clone ID No:Z") provides the unique clone identifier for each clone as previously described and indicated in Tables 1A, 1B, 1C, and 1D. The third column ("AA SEQ ID NO:Y") indicates the Sequence Listing SEQ ID Number for polypeptide sequences encoded by the corresponding cDNA clones (also as indicated in Tables 1A, 1B, and 2). The fourth column ("Biological Activity") indicates a biological activity corresponding to the indicated polypeptides (or polynucleotides encoding said polypeptides). The fifth column ("Exemplary Activity Assay") further describes the corresponding biological activity and also provides information pertaining to the various types of assays which may be performed to test, demonstrate, or quantify the corresponding biological activity. The sixth column ("Preferred Indictions") describes particular embodiments of the invention as well as indications (e.g. pathologies, diseases, disorders, abnormalities, etc.) for which polynucleotides and polypeptides of the invention (including antibodies, agonists, and/or antagonists thereof) may be used in detecting, diagnosing, preventing, and/or treating.

[121] Table 1E describes the use of, inter alia, FMAT technology for testing or demonstrating various biological activities. Fluorometric microvolume assay technology (FMAT) is a fluorescence-based system which provides a means to perform nonradioactive cell- and bead-based assays to detect activation of cell signal transduction pathways. This technology was designed specifically for ligand binding and immunological assays. Using this technology, fluorescent cells or beads at the bottom of the well are detected as localized areas of concentrated fluorescence using a data processing system. Unbound flurophore comprising the background signal is ignored,

allowing for a wide variety of homogeneous assays. FMAT technology may be used for peptide ligand binding assays, immunofluorescence, apoptosis, cytotoxicity, and bead-based immunocapture assays. See, Miraglia S et. al., "Homogeneous cell and bead based assays for highthroughput screening using flourometric microvolume assay technology," Journal of Biomolecular Screening; 4:193-204 (1999). In particular, FMAT technology may be used to test, confirm, and/or identify the ability of polypeptides (including polypeptide fragments and variants) to activate signal transduction pathways. For example, FMAT technology may be used to test, confirm, and/or identify the ability of polypeptides to upregulate production of immunomodulatory proteins (such as, for example, interleukins, GM-CSF, Rantes, and Tumor Necrosis factors, as well as other cellular regulators (e.g. insulin)).

[122] Table 1E also describes the use of kinase assays for testing, demonstrating, or quantifying biological activity. In this regard, the phosphorylation and de-phosphorylation of specific amino acid residues (e.g. Tyrosine, Serine, Threonine) on cell-signal transduction proteins provides a fast, reversible means for activation and de-activation of cellular signal transduction pathways. Moreover, cell signal transduction via phosphorylation/de-phosphorylation is crucial to the regulation of a wide variety of cellular processes (e.g. proliferation, differentiation, migration, apoptosis, etc.). Accordingly, kinase assays provide a powerful tool useful for testing, confirming, and/or identifying polypeptides (including polypeptide fragments and variants) that mediate cell signal transduction events via protein phosphorylation. See e.g., Forrer, P., Tamaskovic R., and Jaussi, R. "Enzyme-Linked Immunosorbent Assay for Measurement of JNK, ERK, and p38 Kinase Activities" Biol. Chem. 379(8-9): 1101-1110 (1998).

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Gene	cDNA	AA SEQ ID	Biological Activity	Exemplary Activity Assay	Preferred Indications
No.	Clone ID	NO: Y			
-	H6BSF56	515	Activation of	Assays for the activation of transcription	A preferred embodiment of the invention includes a
			transcription through	through the Serum Response Element	method for inhibiting (e.g., reducing) TNF alpha
			serum response element	(SRE) are well-known in the art and may	production. An alternative preferred embodiment of the
			in immune cells (such	be used or routinely modified to assess the	invention includes a method for stimulating (e.g.,
			as T-cells).	ability of polypeptides of the invention	increasing) TNF alpha production. Preferred indications
				(including antibodies and agonists or	include blood disorders (e.g., as described below under
				antagonists of the invention) to regulate	"Immune Activity", "Blood-Related Disorders", and/or
				the serum response factors and modulate	"Cardiovascular Disorders"), Highly preferred indications
				the expression of genes involved in	include autoimmune diseases (e.g., rheumatoid arthritis,
				growth. Exemplary assays for	systemic lupus erythematosis, Crohn's disease, multiple
				transcription through the SRE that may be	sclerosis and/or as described below), immunodeficiencies
				used or routinely modified to test SRE	(e.g., as described below), boosting a T cell-mediated
				activity of the polypeptides of the	immune response, and suppressing a T cell-mediated
				invention (including antibodies and	immune response. Additional highly preferred indications
				agonists or antagonists of the invention)	include inflammation and inflammatory disorders, and
				include assays disclosed in Berger et al.,	treating joint damage in patients with rheumatoid arthritis.
				Gene 66:1-10 (1998); Cullen and Malm,	An additional highly preferred indication is sepsis.
				Methods in Enzymol 216:362-368 (1992);	Highly preferred indications include neoplastic diseases
				Henthorn et al., Proc Natl Acad Sci USA	(e.g., leukemia, lymphoma, and/or as described below
				85:6342-6346 (1988); and Black et al.,	under "Hyperproliferative Disorders"). Additionally,
				Virus Genes 12(2):105-117 (1997), the	highly preferred indications include neoplasms and
				content of each of which are herein	cancers, such as, for example, leukemia, lymphoma,
				incorporated by reference in its entirety. T	melanoma, glioma (e.g., malignant glioma), solid tumors,
				cells that may be used according to these	and prostate, breast, lung, colon, pancreatic, esophageal,
				assays are publicly available (e.g., through	stomach, brain, liver and urinary cancer. Other preferred
				the ATCC). Exemplary mouse T cells that	indications include benign dysproliferative disorders and
				may be used according to these assays	pre-neoplastic conditions, such as, for example,
				include the CTLL cell line, which is an IL-	hyperplasia, metaplasia, and/or dysplasia. Preferred
				2 dependent suspension culture of T cells	indications include anemia, pancytopenia, leukopenia,
				with cytotoxic activity.	thrombocytopenia, Hodgkin's disease, acute lymphocytic

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					anemia (ALL), plasmacytomas, multiple myeloma, Burkitt's lymphoma, arthritis, AIDS, granulomatous disease, inflammatory bowel disease, neutropenia, neutrophilia, psoriasis, suppression of immune reactions to transplanted organs and tissues, hemophilia, hypercoagulation, diabetes mellitus, endocarditis, meningitis, Lyme Disease, cardiac reperfusion injury, and asthma and allergy. An additional preferred indication is infection (e.g., an infectious disease as described below under "Infectious Disease").
2	Н6ЕDM64	516	Insulin Secretion	Assays for measuring secretion of insulin are well-known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to stimulate insulin secretion. For example, insulin secretion is measured by FMAT using anti-rat insulin antibodies. Insulin secretion from pancreatic beta cells is upregulated by glucose and also by certain proteins/peptides, and disregulation is a key component in diabetes. Exemplary assays that may be used or routinely modified to test for stimulation of insulin secretion (from pancreatic cells) by polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in: Shimizu, H., et al., Endocr J, 47(3):261-9 (2000); Salapatek, A.M., et al., Mol Endocrinol, 13(8):1305-17 (1999); Filipsson, K., et al., Ann N Y Acad Sci, 865:441-4 (1998); Olson, L.K., et al., J Biol Chem, 271(28):16544-52 (1996); and, Miraglia S et al., Journal of Biomolecular	A highly preferred indication is diabetes mellitus. An additional highly preferred indication is a complication associated with diabetes (e.g., diabetic retinopathy, diabetes (e.g., diabetic retinopathy, diabetes) diabetic nephropathy, kidney disease (e.g., renal failure, nephropathy and/or other diseases and disorders as described in the "Renal Disorders" section below), diabetic neuropathy, nerve disease and nerve damage (e.g., due to diabetic neuropathy, nerve disease and nerve damage (e.g., due to diabetic neuropathy or blood vessel blockage), seizures, mental confusion, drowsiness, nonketotic hyperglycemic-hyperosmolar coma, cardiovascular disease (e.g., heart disease, atherosclerosis, microvascular disease, hypertension, stroke, and other diseases and disorders as described in the "Endocrine Disorders" section below), neuropathy, vision impairment (e.g., diabetic retinopathy and blindness), ulcers and impaired wound healing, and infection (e.g., infectious Diseases and disorders as described in the "Infectious Diseases" section below, especially of the urinary tract and skin), carpal tunnel syndrome and Dupuytren's contracture).  An additional highly preferred indications include weight loss or alternatively, weight gain.  Additional highly preferred indications include weight loss or alternatively, weight gain.

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with insulin resistance.	A preferred embodiment of the invention includes a method for inhibiting (e.g., reducing) TNF alpha production. An alternative highly preferred embodiment of production. An alternative highly preferred embodiment of increasing) TNF alpha production. Preferred indications include blood disorders (e.g., as described below under "Tmmune Activity", "Blood-Related Disorders", and/or "Cardiovascular Disorders"), Highly preferred indications include autoimmune diseases (e.g., rheumatoid arthritis, systemic lupus erythematosis, Crohn's disease, multiple systemic lupus erythematosis, Crohn's disease, multiple systemic lupus erythematosis, Crohn's disease, multiple systemic lupus erythematosis, An delicional highly preferred indications include inflammation and inflammatory disorders, and include inflammation and inflammatory disorders, and treating joint damage in patients with rheumatoid arthritis. An additional highly preferred indication is sepsis. Highly preferred indications include neoplastic diseases
orated by tic cells see assays ligh the generated. may be nolude disherent om Syrian with SV40. I receptors. is gon and 1777. Iem. J. 219: Vatl. Acad.	Sci. USA 78: 4339-4343, 1981. Assays for the activation of transcription through the Serum Response Element (SRE) are well-known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to regulate serum response factors and modulate the expression of genes involved in growth and upregulate the function of growthrelated genes in many cell types.  Exemplary assays for transcription through the SRE that may be used or routinely modified to test SRE activity of the polypeptides of the invention (including antibodies and agonists or antagonists of antibodies and agonists or antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen
	Activation of transcription through serum response element in immune cells (such as natural killer cells).
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(e.g., leukemia, lymphoma, and/or as described below under "Hyperproliferative Disorders"). Additionally, highly preferred indications include neoplasms and cancers, such as, for example, leukemia, lymphoma, melanoma, glioma (e.g., malignant glioma), solid tumors, and prostate, breast, lung, colon, pancreatic, esophageal, stomach, brain, liver and urinary cancer. Other preferred indications include benign dysproliferative disorders and pre-neoplastic conditions, such as, for example, hyperplasia, metaplasia, and/or dysplasia. Preferred indications include anemia, pancytopenia, leukopenia, ethrombocytopenia, Hodgkin's disease, acute lymphocytic anemia (ALL), plasmacytomas, multiple myeloma, Burkitt's lymphoma, arthritis, AIDS, granulomatous disease, inflammatory bowel disease, neutropenia, neutrophilia, psoriasis, suppression of immune reactions to transplanted organs and tissues, hemophilia, hypercoagulation, diabetes mellitus, endocarditis, meningitis, Lyme Disease, cardiac reperfusion injury, and asthma and allergy. An additional preferred indication is infection (e.g., an infectious disease as described below under "Infectious Disease").	Preferred indications include blood disorders (e.g., as described below under "Immune Activity", "Blood-Related Disorders", and/or "Cardiovascular Disorders"), and infection (e.g., an infectious disease as described below under "Infectious Disease"). Preferred indications include autoimmune diseases (e.g., rheumatoid arthritis, systemic lupus erythematosis, multiple sclerosis and/or as described below), immunodeficiencies (e.g., as described below), boosting a T cell-mediated immune response. Additional preferred indications include inflammation and inflammatory disorders. Highly preferred indications include neoplastic diseases (e.g., leukemia, lymphoma, and/or as described below under "Hyperproliferative Disorders"). Highly preferred
and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Benson et al., J Immunol 153(9):3862-3873 (1994); and Black et al., Virus Genes 12(2):105-117 (1997), the content of each of which are herein incorporated by reference in its entirety. T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the NK-YT cell line, which is a human natural killer cell line with cytolytic and cytotoxic activity.	Assays for the activation of transcription through the cAMP response element are well-known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to increase cAMP and regulate CREB transcription factors, and modulate expression of genes involved in a wide variety of cell functions. Exemplary assays for transcription through the cAMP response element that may be used or routinely modified to test cAMP-response element activity of polypeptides of the invention (including antibodies and
	Activation of transcription through cAMP response element in immune cells (such as T-cells).
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agonists or antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Black et al., Virus Genes 15(2):105-117 (1997); and Belkowski et al., J Immunol 161(2):659- 665 (1998), the contents of each of which are herein incorporated by reference in its entirety. T cells that may be used according to these assays are publicly available (e.g., through the ATCC).  Exemplary mouse T cells that may be used according to these assays include the CTLL cell line, which is a suspension culture of IL-2 dependent cytotoxic T me	Activation of the activation of transcription transcription through the Serum Response Element serum response element (SRE) are well-known in the art and may in immune cells (such a be used or routinely modified to assess the arrangement cells (such a serum response element as T-cells).  Appeterred embodiment of the invention in immune cells (such a be used or routinely modified to assess the in immune cells (such a serum response factors and agonists of the invention) to regulate expression of genes involved in growth. Exemplary assays for used or routinely modified to test SRE activity of the polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992);  Apprendic of the invention includes a method for inhibiting (e.g., reducing) TNF alpha production. An alternative preferred embodiment of the invention includes as and or an argonists of the invention include assays disclosed in Berger et al., immune response. Additional highly preferred indications include assays disclosed in Berger et al., immune response indications include assays disclosed in Berger et al., immune response indications include assays disclosed in Berger et al., immune response indications include assays disclosed in Berger et al., immune response indications include assays disclosed in Berger et al., immune response indications is sepsis.
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				85:6342-6346 (1988); and Black et al., Virus Genes 12(2):105-117 (1997), the content of each of which are herein incorporated by reference in its entirety. T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary mouse T cells that may be used according to these assays include the CTLL cell line, which is an IL-2 dependent suspension culture of T cells with cytotoxic activity.	under "Hyperproliferative Disorders"). Additionally, highly preferred indications include neoplasms and cancers, such as, for example, leukemia, lymphoma, melanoma, glioma (e.g., malignant glioma), solid tumors, and prostate, breast, lung, colon, pancreatic, esophageal, stomach, brain, liver and urinary cancer. Other preferred indications include benign dysproliferative disorders and pre-neoplastic conditions, such as, for example, hyperplasia, metaplasia, and/or dysplasia. Preferred indications include anemia, pancytopenia, leukopenia, thrombocytopenia, Hodgkin's disease, acute lymphocytic anemia (ALL), plasmacytomas, multiple myeloma, Burkitt's lymphoma, arthritis, AIDS, granulomatous disease, inflammatory bowel disease, neutropenia, neutrophilia, psoriasis, suppression of immune reactions to transplanted organs and tissues, hemophilia, mortansplanted organs and tissues, hemophilia, mortansplanted organs and tissues, hemophilia, and asthma and allergy. An additional preferred indication is infection (e.g., an infectious disease as described below under "Infectious Disease").
4	HACAB68	518	Activation of Endothelial Cell p38 or JNK Signaling Pathway.	Kinase assay. INK and p38 kinase assays for signal transduction that regulate cell proliferation, activation, or apoptosis are well known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to promote or inhibit cell proliferation, activation, and apoptosis. Exemplary assays for JNK and p38 kinase activity that may be used or routinely modified to test JNK and p38 kinase induced activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include the assays disclosed in Forrer et	A highly preferred embodiment of the invention includes a method for stimulating endothelial cell growth. An alternative highly preferred embodiment of the invention includes a method for inhibiting endothelial cell growth. A highly preferred embodiment of the invention includes a method for stimulating endothelial cell proliferation. An alternative highly preferred embodiment of the invention includes a method for inhibiting endothelial cell proliferation. A highly preferred embodiment of the invention includes a method for stimulating apoptosis of endothelial cells. An alternative highly preferred embodiment of the invention includes a method for inhibiting (e.g., decreasing) apoptosis of endothelial cells. A highly preferred embodiment of the invention includes a method for stimulating (e.g., increasing) endothelial cell activation.

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al., Biol Chem 379(8-9):1101-1110 (1998); Gupta et al., Exp Cell Res 247(2):	An alternative highly preferred embodiment of the invention includes a method for inhibiting (e.g.,
495-504 (1999); Kyriakis JM, Biochem Soc Symp 64:29-48 (1999); Chang and	decreasing) the activation of anglor inactivating endothelial cells.  A highly preferred embodiment of
Karin, Nature 410(6824):37-40 (2001);	the invention includes a method for stimulating
and Cobb MH, Prog Biophys Mol Biol	angiogenisis. An alternative highly preferred embodiment
71(3-4):479-500 (1999); the contents of	of the invention includes a method for inhibiting
each of which are never incorporated by reference in its entirety. Endothelial cells	angiogenesis. A nignij prejerred embodiment of the invention includes a method for reducing cardiac
that may be used according to these assays	hypertrophy. An alternative highly preferred embodiment
are publicly available (e.g., through the	of the invention includes a method for inducing cardiac
ATCC). Exemplary endothelial cells that	hypertrophy. Highly preferred indications include
may be used according to these assays	neoplastic diseases (e.g., as described below under
include human umbilical vein endothelial	"Hyperproliferative Disorders"), and disorders of the
cells (HUVEC), which are endothelial	cardiovascular system (e.g., heart disease, congestive heart
 cells which line venous blood vessels, and	failure, hypertension, aortic stenosis, cardiomyopathy,
are involved in functions that include, but	valvular regurgitation, left ventricular dysfunction,
are not limited to, angiogenesis, vascular	atherosclerosis and atherosclerotic vascular disease,
permeability, vascular tone, and immune	diabetic nephropathy, intracardiac shunt, cardiac
cell extravasation.	hypertrophy, myocardial infarction, chronic hemodynamic
	overload, and/or as described below under
	"Cardiovascular Disorders"). Highly preferred indications
	include cardiovascular, endothelial and/or angiogenic
	disorders (e.g., systemic disorders that affect vessels such
	as diabetes mellitus, as well as diseases of the vessels
	themselves, such as of the arteries, capillaries, veins and/or
	lymphatics). Highly preferred are indications that
	stimulate angiogenesis and/or cardiovascularization.
	Highly preferred are indications that inhibit angiogenesis
	and/or cardiovascularization. Highly preferred
	indications include antiangiogenic activity to treat solid
	tumors, leukemias, and Kaposi's sarcoma, and retinal
	disorders. Highly preferred indications include neoplasms
	and cancer, such as, Kaposi's sarcoma, hemangioma
	(capillary and cavernous), glomus tumors, telangiectasia,
	bacillary angiomatosis, hemangioendothelioma,
	angiosarcoma, naemangiopericytoma, iympnangioma,

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					and Crohn's disease), and pain management.
4	HACAB68	518	Activation of Skeletal	Kinase assay. Kinase assays, for example	A highly preferred embodiment of the invention
			Mucle Cell PI3 Kinase	an GSK-3 kinase assay, for PI3 kinase	includes a method for increasing muscle cell survival An
			Signalling Pathway	signal transduction that regulate glucose	alternative highly preferred embodiment of the invention
				metabolism and cell survivial are well-	includes a method for decreasing muscle cell survival.
				known in the art and may be used or	A preferred embodiment of the invention includes a
				routinely modified to assess the ability of	method for stimulating muscle cell proliferation. In a
				polypeptides of the invention (including	specific embodiment, skeletal muscle cell proliferation is
				antibodies and agonists or antagonists of	stimulated. An alternative highly preferred embodiment of
				the invention) to promote or inhibit	the invention includes a method for inhibiting muscle cell
				glucose metabolism and cell survival.	proliferation. In a specific embodiment, skeletal muscle
		·· ,		Exemplary assays for PI3 kinase activity	cell proliferation is inhibited. A preferred embodiment
				that may be used or routinely modified to	of the invention includes a method for stimulating muscle
				test PI3 kinase-induced activity of	cell differentiation. In a specific embodiment, skeletal
				polypeptides of the invention (including	muscle cell differentiation is stimulated. An alternative
				antibodies and agonists or antagonists of	highly preferred embodiment of the invention includes a
				the invention) include assays disclosed in	method for inhibiting muscle cell differentiation. In a
				Forrer et al., Biol Chem 379(8-9):1101-	specific embodiment, skeletal muscle cell differentiation is
				1110 (1998); Nikoulina et al., Diabetes	inhibited. Highly preferred indications include disorders
				49(2):263-271 (2000); and Schreyer et al.,	of the musculoskeletal system. Preferred indications
				Diabetes 48(8):1662-1666 (1999), the	include neoplastic diseases (e.g., as described below under
				contents of each of which are herein	"Hyperproliferative Disorders"), endocrine disorders (e.g.,
				incorporated by reference in its entirety.	as described below under "Endocrine Disorders"), neural
				Rat myoblast cells that may be used	disorders (e.g., as described below under "Neural Activity
				according to these assays are publicly	and Neurological Diseases"), blood disorders (e.g., as
				available (e.g., through the ATCC).	described below under "Immune Activity",
				Exemplary rat myoblast cells that may be	"Cardiovascular Disorders", and/or "Blood-Related
				used according to these assays include L6	Disorders"), immune disorders (e.g., as described below
				cells. L6 is an adherent rat myoblast cell	under "Immune Activity"), and infection (e.g., as
				line, isolated from primary cultures of rat	described below under "Infectious Disease"). A
	•			thigh muscle, that fuses to form	highly preferred indication is diabetes mellitus. An
				multinucleated myotubes and striated	additional highly preferred indication is a complication
				fibers after culture in differentiation media.	associated with diabetes (e.g., diabetic retinopathy,
					diabetic nephropathy, kidney disease (e.g., renal failure,
					nephropathy and/or other diseases and disorders as
					described in the "Renal Disorders" section below), diabetic
					neuropathy, nerve disease and nerve damage (e.g., due to

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					diabetic neuropathy), blood vessel blockage, heart disease,
					stroke, impotence (e.g., due to diabetic neuropathy or
					blood vessel blockage), seizures, mental confusion,
					drowsiness, nonketotic hyperglycemic-hyperosmolar
					coma, cardiovascular disease (e.g., heart disease,
					atherosclerosis, microvascular disease, hypertension,
					stroke, and other diseases and disorders as described in the
					"Cardiovascular Disorders" section below), dyslipidemia,
					endocrine disorders (as described in the "Endocrine
					Disorders" section below), neuropathy, vision impairment
					(e.g., diabetic retinopathy and blindness), ulcers and
					impaired wound healing, infections (e.g., infectious
					diseases and disorders as described in the "Infectious
					Diseases" section below, especially of the urinary tract and
					skin), carpal tunnel syndrome and Dupuytren's
					contracture). An additional highly preferred indication
					is obesity and/or complications associated with obesity.
					Additional highly preferred indications include weight loss
<b></b>					or alternatively, weight gain. Additional highly
	•				preferred indications are complications associated with
					insulin resistance. Additonal highly preferred
					indications are disorders of the musculoskeletal system
					including myopathies, muscular dystrophy, and/or as
					described herein. Additional highly preferred
					indications include: myopathy, atrophy, congestive heart
					failure, cachexia, myxomas, fibromas, congenital
					e, c
			-		heart valve disease, and vascular disease. Highly
					preferred indications include neoplasms and cancer, such
					as, rhabdomyoma, rhabdosarcoma, stomach, esophageal,
					prostate, and urinary cancer. Preferred indications also
					include breast, lung, colon, pancreatic, brain, and liver
					cancer. Other preferred indications include benign
					dysproliferative disorders and pre-neoplastic conditions,
					such as, hyperplasia, metaplasia, and/or dysplasia.
~	HACBJ56	519		Assays for the regulation of viability and	A highly preferred indication is diabetes mellitus.
			and proliferation of	proliferation of cells in vitro are well-	An additional highly preferred indication is a complication

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	pancrea	pancreatic beta cells.	known in the art and may be used or	associated with diabetes (e.g., diabetic retinopathy,
			routinely modified to assess the ability of	diabetic nephropathy, kidney disease (e.g., renal failure,
			polypeptides of the invention (including	nephropathy and/or other diseases and disorders as
			antibodies and agonists or antagonists of	described in the "Renal Disorders" section below), diabetic
-			the invention) to regulate viability and	neuropathy, nerve disease and nerve damage (e.g., due to
			proliferation of pancreatic beta cells. For	diabetic neuropathy), blood vessel blockage, heart disease,
			example, the Cell Titer-Glo luminescent	stroke, impotence (e.g., due to diabetic neuropathy or
			cell viability assay measures the number of	blood vessel blockage), seizures, mental confusion,
			viable cells in culture based on	drowsiness, nonketotic hyperglycemic-hyperosmolar
			quantitation of the ATP present which	coma, cardiovascular disease (e.g., heart disease,
			signals the presence of metabolically	atherosclerosis, microvascular disease, hypertension,
			active cells. Exemplary assays that may be	stroke, and other diseases and disorders as described in the
			used or routinely modified to test	"Cardiovascular Disorders" section below), dyslipidemia,
			regulation of viability and proliferation of	endocrine disorders (as described in the "Endocrine
			pancreatic beta cells by polypeptides of the	Disorders" section below), neuropathy, vision impairment
			invention (including antibodies and	(e.g., diabetic retinopathy and blindness), ulcers and
			agonists or antagonists of the invention)	impaired wound healing, and infection (e.g., infectious
			include assays disclosed in: Friedrichsen	diseases and disorders as described in the "Infectious
			BN, et al., Mol Endocrinol, 15(1):136-48	Diseases" section below, especially of the urinary tract and
			(2001); Huotari MA, et al., Endocrinology,	skin), carpal tunnel syndrome and Dupuytren's
			139(4):1494-9 (1998); Hugl SR, et al., J	contracture). An additional highly preferred
			Biol Chem 1998 Jul 10;273(28):17771-9	indication is obesity and/or complications associated with
			(1998), the contents of each of which is	obesity. Additional highly preferred indications include
			herein incorporated by reference in its	weight loss or alternatively, weight gain. Aditional
			entirety. Pancreatic cells that may be used	highly preferred indications are complications associated
			according to these assays are publicly	with insulin resistance.
			available (e.g., through the ATCC) and/or	
			may be routinely generated. Exemplary	
			pancreatic cells that may be used	
			according to these assays include rat INS-1	
			cells. INS-1 cells are a semi-adherent cell	
			line established from cells isolated from an	
	-		X-ray induced rat transplantable	
			insulinoma. These cells retain	
			characteristics typical of native pancreatic	
			beta cells including glucose inducible	
			insulin secretion. References: Asfari et al.	

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$\vdash$				Endocrinology 1992 130:167.	
T	HACBS22	520	Production of ICAM-1	Assays for measuring expression of	Preferred embodiments of the invention include using
		)		ICAM-1 are well-known in the art and	polypeptides of the invention (or antibodies, agonists, or
				may be used or routinely modified to	antagonists thereof) in detection, diagnosis, prevention,
				assess the ability of polypeptides of the	and/or treatment of Vascular Disease, Atherosclerosis,
				invention (including antibodies and	Restenosis, Stroke, and Asthma.
				agonists or antagonists of the invention) to	
				regulate ICAM-1 expression. Exemplary	
				assays that may be used or routinely	
				modified to measure ICAM-1 expression	
				include assays disclosed in: Rolfe BE, et	
				al., Atherosclerosis, 149(1):99-110 (2000);	
				Panettieri RA Jr, et al., J Immunol,	
				154(5):2358-2365 (1995); and, Grunstein	
				MM, et al., Am J Physiol Lung Cell Mol	
				Physiol, 278(6):L1154-L1163 (2000), the	
				contents of each of which is herein	
				incorporated by reference in its entirety.	
				Cells that may be used according to these	
				assays are publicly available (e.g., through	
				the ATCC) and/or may be routinely	
				generated. Exemplary cells that may be	
				used according to these assays include	
				Aortic Smooth Muscle Cells (AOSMC);	
				such as bovine AOSMC.	
	HADDE71	521	Activation of	Assays for the activation of transcription	A highly preferred indication is allergy. Another
			transcription through	through the Signal Transducers and	highly preferred indication is asthma. Additional
			STAT6 response	Activators of Transcription (STAT6)	highly preferred indications include inflammation and
			element in immune	response element are well-known in the art	inflammatory disorders. Preferred indications
			cells (such as natural	and may be used or routinely modified to	include blood disorders (e.g., as described below under
			killer cells).	assess the ability of polypeptides of the	"Immune Activity", "Blood-Related Disorders", and/or
				invention (including antibodies and	"Cardiovascular Disorders"). Preferred indications include
				agonists or antagonists of the invention) to	autoimmune diseases (e.g., rheumatoid arthritis, systemic
				regulate STAT6 transcription factors and	lupus erythematosis, multiple sclerosis and/or as described
				modulate the expression of multiple genes.	below) and immunodeficiencies (e.g., as described below).
				Exemplary assays for transcription through	Preferred indications include neoplastic diseases (e.g.,
7				the STATO response element that may be	leukemia, lymphoma, melanoma, and/or as described

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				used or routinely modified to test STAT6	below under "Hyperproliferative Disorders"). Preferred
				response element activity of the	indications include neoplasms, such as, for example,
				polypeptides of the invention (including	leukemia, lymphoma, melanoma, and prostate, breast,
				antibodies and agonists or antagonists of	lung, colon, pancreatic, esophageal, stomach, brain, liver
				the invention) include assays disclosed in	and urinary cancer. Other preferred indications include
				Berger et al., Gene 66:1-10 (1998); Cullen	benign dysproliferative disorders and pre-neoplastic
				and Malm, Methods in Enzymol 216:362-	as, for examp
				368 (1992); Henthorn et al., Proc Natl	and/or dysplasia. Preferred indications include
				Acad Sci USA 85:6342-6346 (1988);	anemia, pancytopenia, leukopenia, thrombocytopenia,
				Georas et al., Blood 92(12):4529-4538	Hodgkin's disease, acute lymphocytic anemia (ALL),
				(1998); Moffatt et al., Transplantation	plasmacytomas, multiple myeloma, Burkitt's lymphoma,
				69(7):1521-1523 (2000); Curiel et al., Eur	arthritis, AIDS, granulomatous disease, inflammatory
				J Immunol 27(8):1982-1987 (1997); and	bowel disease, sepsis, neutropenia, neutrophilia, psoriasis,
				Masuda et al., J Biol Chem	suppression of immune reactions to transplanted organs
				275(38):29331-29337 (2000), the contents	and tissues, hemophilia, hypercoagulation, diabetes
				of each of which are herein incorporated	mellitus, endocarditis, meningitis, and Lyme Disease.
				by reference in its entirety. T cells that	Additional preferred indications include infection (e.g., an
				may be used according to these assays are	infectious disease as described below under "Infectious
				publicly available (e.g., through the	Disease").
				ATCC). Exemplary rat natural killer cells	
				that may be used according to these assays	
				are publicly available (e.g., through the	
				ATCC).	
8	HADDJ13	522	Activation of	Assays for the activation of transcription	A highly preferred indication is allergy. Another
			transcription through	through the Signal Transducers and	highly preferred indication is asthma. Additional
			STAT6 response	Activators of Transcription (STAT6)	highly preferred indications include inflammation and
			element in immune	response element are well-known in the art	inflammatory disorders. Preferred indications
			cells (such as natural	and may be used or routinely modified to	include blood disorders (e.g., as described below under
			killer cells).	assess the ability of polypeptides of the	"Immune Activity", "Blood-Related Disorders", and/or
				invention (including antibodies and	"Cardiovascular Disorders"). Preferred indications include
				agonists or antagonists of the invention) to	autoimmune diseases (e.g., rheumatoid arthritis, systemic
				regulate STAT6 transcription factors and	lupus erythematosis, multiple sclerosis and/or as described
				modulate the expression of multiple genes.	below) and immunodeficiencies (e.g., as described below).
				Exemplary assays for transcription through	Preferred indications include neoplastic diseases (e.g.,
				the STAT6 response element that may be	leukemia, lymphoma, melanoma, and/or as described
				used or routinely modified to test STAT6	below under "Hyperproliferative Disorders"). Preferred
				response element activity of the	indications include neoplasms, such as, for example,

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				polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Georas et al., Blood 92(12):4529-4538	leukemia, lymphoma, melanoma, and prostate, breast, lung, colon, pancreatic, esophageal, stomach, brain, liver and urinary cancer. Other preferred indications include benign dysproliferative disorders and pre-neoplastic conditions, such as, for example, hyperplasia, metaplasia, and/or dysplasia.  Preferred indications include anemia, pancytopenia, leukopenia, thrombocytopenia, Hodekin's disease, acute lymphocytic anemia (ALL).
			·	(1998); Moffatt et al., Transplantation 69(7):1521-1523 (2000); Curiel et al., Eur J Immunol 27(8):1982-1987 (1997); and Masuda et al., J Biol Chem 275(38):29331-29337 (2000), the contents of each of which are herein incorporated	plasmacytomas, multiple myeloma, Burkitt's lymphoma, arthritis, AIDS, granulomatous disease, inflammatory bowel disease, sepsis, neutropenia, neutrophilia, psoriasis, suppression of immune reactions to transplanted organs and tissues, hemophilia, hypercoagulation, diabetes mellitus, endocarditis, meningits, and I yme Disease
				by reference in its entirety. T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary rat natural killer cells that may be used according to these assays are publicly available (e.g., through the	Additional preferred indications include infection (e.g., an infectious disease as described below under "Infectious Disease").
∞	HADDJ13	522	Activation of transcription through GAS response element in immune cells (such as T-cells).	ASSAYS for the activation of transcription through the Gamma Interferon Activation Site (GAS) response element are well-known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to regulate STAT transcription factors and modulate gene expression involved in a wide variety of	Highly preferred indications include neoplastic diseases (e.g., leukemia, lymphoma, and/or as described below under "Hyperproliferative Disorders"). Highly preferred indications include neoplasms and cancers, such as, for example, leukemia, lymphoma (e.g., T cell lymphoma, Burkitt's lymphoma, non-Hodgkins lymphoma, Hodgkin's disease), melanoma, and prostate, breast, lung, colon, pancreatic, esophageal, stomach, brain, liver and urinary cancer. Other preferred indications include benign dysproliferative disorders and pre-neoplastic conditions.
				cell functions. Exemplary assays for transcription through the GAS response element that may be used or routinely modified to test GAS-response element activity of polypeptides of the invention (including antibodies and agonists or	such as, for example, hyperplasia, metaplasia, and/or dysplasia. Preferred indications include autoimmune diseases (e.g., rheumatoid arthritis, systemic lupus erythematosis, multiple sclerosis and/or as described below), immunodeficiencies (e.g., as described below), boosting a T cell-mediated immune response, and

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referred indications ribed below under Disorders", and/or ection (e.g., viral sociated with chronic at osteoporosis, bed below under preferred indication Preferred indications benia, anemia (ALL), rthritis, AIDS, bowel disease, riasis, suppression of ans and tissues, es mellitus, se, and asthma and	ood disorders Activity", iovascular include thritis, systemic d/or as described ribed below), nse, and sponse. clude . An additional g., an infectious ous Disease"). iseases (e.g., below under d indications or example.
suppressing a T cell-mediated immune response.  Additional preferred indications include inflammation and inflammatory disorders. Highly preferred indications include blood disorders (e.g., as described below under "Immune Activity", "Blood-Related Disorders", and/or "Cardiovascular Disorders"), and infection (e.g., viral infections, tuberculosis, infections associated with chronic granulomatosus disease and malignant osteoporosis, and/or an infectious disease as described below under "Infectious Disease"). An additional preferred indication is idiopathic pulmonary fibrosis. Preferred indications include anemia, pancytopenia, leukopenia, thrombocytopenia, acute lymphocytic anemia (ALL), plasmacytomas, multiple myeloma, arthritis, AIDS, granulomatous disease, inflammatory bowel disease, sepsis, neutropenia, neutrophilia, psoriasis, suppression of immune reactions to transplanted organs and tissues, hemophilia, hypercoagulation, diabetes mellitus, endocarditis, meningitis, Lyme Disease, and asthma and allergy.	Highly preferred indications include blood disorders (e.g., as described below under "Immune Activity", "Blood-Related Disorders", and/or "Cardiovascular Disorders"). Highly preferred indications include autoimmune diseases (e.g., rheumatoid arthritis, systemic lupus erythematosis, multiple sclerosis and/or as described below), immunodeficiencies (e.g., as described below), boosting a T cell-mediated immune response, and suppressing a T cell-mediated immune response. Additional highly preferred indications include inflammation and inflammatory disorders. An additional highly preferred indication is infectious Disease"). Preferred indications include neoplastic diseases (e.g., leukemia, lymphoma, and/or as described below under "Hyperproliferative Disorders"). Preferred indications include neoplasms and cancers, such as, for example.
suppressing a 1 Additional prel inflammatory of include blood of "Immune Actival of a cardiovascula infections, tube granulomatosu, and/or an infections Dis is idiopathic puinclude anemia thrombocytope plasmacytomas granulomatous sepsis, neutrop immune reactic hemophilia, hy endocarditis, m allergy.	Highly prefe  (e.g., as descrit "Blood-Related Disorders"). His autoimmune dis lupus erythema below), immun boosting a T ce suppressing a T Additional high inflammation as highly preferred disease as described belowers and the light of the leave indicate indicate leave indicate leave include neoplas include neoplas include neoplas
tion) include ger et al., Gene and Malm, 6:362-368 (1992); atl Acad Sci USA Matikainen et al., (1999); and mol 155(10):4582- its of each of which by reference in its man T cells, such at may be used s are publicly the ATCC).	n of transcription stor of Activated T element are well- y be used or sess the ability of antion (including or antagonists of e NFAT I modulate blved in ctions. Exemplary through the NFAT ay be used or st NFAT-response peptides of the ibodies and of the invention)
antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Matikainen et al., Blood 93(6):1980-1991 (1999); and Henttinen et al., J Immunol 155(10):4582-4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary human T cells, such as the SUPT cell line, that may be used according to these assays are publicly available (e.g., through the ATCC).	Assays for the activation of transcription through the Nuclear Factor of Activated T cells (NFAT) response element are well-known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to regulate NFAT transcription factors and modulate expression of genes involved in immunomodulatory functions. Exemplary assays for transcription through the NFAT response element that may be used or routinely modified to test NFAT-response element activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention)
	Activation of transcription through NFAT response element in immune cells (such as natural killer cells).
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				include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Aramburu et al., J Exp Med 182(3):801-810 (1995); De Boer et al., Int J Biochem Cell Biol 31(10):1221-1236 (1999); Fraser et al., Eur J Immunol 29(3):838-844 (1999); and Yeseen et al., J Biol Chem 268(19):14285-14293 (1993), the contents of each of which are herein incorporated by reference in its entirety. NK cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary human NK cells that may be used according to these assays include the NK-YT cell line, which is a human natural killer cell line with cytolytic and cytotoxic activity.	leukemia, lymphoma, and prostate, breast, lung, colon, pancreatic, esophageal, stomach, brain, liver and urinary cancer. Other preferred indications include benign dysproliferative disorders and pre-neoplastic conditions, such as, for example, hyperplasia, metaplasia, and/or dysplasia. Preferred indications also include anemia, pancytopenia, leukopenia, thrombocytopenia, Hodgkin's disease, acute lymphocytic anemia (ALL), plasmacytomas, multiple myeloma, Burkitt's lymphoma, arthritis, AIDS, granulomatous disease, inflammatory bowel disease, sepsis, neutropenia, neutrophilia, psoriasis, suppression of immune reactions to transplanted organs and tissues, hemophilia, hypercoagulation, diabetes mellitus, endocarditis, meningitis, Lyme Disease, asthma and allergy.
6	HADMB15	523	Regulation of transcription via DMEF1 response element in adipocytes and pre-adipocytes	Assays for the regulation of transcription through the DMEF1 response element are well-known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to activate the DMEF1 response element in a reporter construct (such as that containing the GLUT4 promoter) and to regulate insulin production. The DMEF1 response element is present in the GLUT4 promoter and binds to MEF2 transcription factor and another transcription factor that is required for insulin regulation of Glut4 expression in skeletal muscle. GLUT4 is the primary insulin-responsive glucose transporter in	A highly preferred indication is diabetes mellitus. An additional highly preferred indication is a complication associated with diabetes (e.g., diabetic retinopathy, diabetic nephropathy, kidney disease (e.g., renal failure, nephropathy and/or other diseases and disorders as described in the "Renal Disorders" section below), diabetic neuropathy, nerve disease and nerve damage (e.g., due to diabetic neuropathy), blood vessel blockage, heart disease, stroke, impotence (e.g., due to diabetic neuropathy or blood vessel blockage), seizures, mental confusion, drowsiness, nonketotic hyperglycemic-hyperosmolar coma, cardiovascular disease (e.g., heart disease, atherosclerosis, microvascular disease, hypertension, stroke, and other diseases and disorders as described in the "Cardiovascular Disorders" section below), dyslipidemia, endocrine disorders (as described in the "Endocrine Disorders" section below), neuropathy, vision impairment

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				fat and muscle tissue. Exemplary assays that may be used or routinely modified to test for DMEF1 response element activity (in adipocytes and pre-adipocytes) by polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in Thai, M.V., et al., J Biol Chem, 273(23):14285-92 (1998); Mora, S., et al., J Biol Chem, 275(21):16323-8 (2000); Liu, M.L., et al., J Biol Chem, 269(45):28514-21 (1994); "Identification of a 30-base pair regulatory element and novel DNA binding protein that regulates the human GLUT4 promoter in transgenic mice", J Biol Chem. 2000 Aug 4;275(31):23666-73; Berger, et al., Gene 66:1-10 (1988); and, Cullen, B., et al., Methods in Enzymol. 216:362–368 (1992), the contents of each of which is herein incorporated by reference in its entirety. Adipocytes and pre-adipocytes that may be used according to these assays include the mouse 3T3-L1 cell line which is an adherent mouse preadipocyte cell line. Mouse 3T3-L1 cells are a continuous substrain of 3T3 fibroblasts developed through clonal isolation. These cells undergo a pre-adipocyte to adinose-	impaired wound healing, and infection (e.g., infectious diseases and disorders as described in the "Infectious diseases and disorders as described in the "Infectious Diseases" section below, especially of the urinary tract and skin), carpal tunnel syndrome and Dupuytren's contracture). An additional highly preferred indication is obesity and/or complications associated with obesity. Additional highly preferred indications include weight loss or alternatively, weight gain. Additional highly preferred indications are complications associated with insulin resistance.
				like conversion under appropriate differentiation culture conditions.	
6	HADMB15	523	Regulation of apoptosis of immune cells (such as mast cells).	Caspase Apoptosis. Assays for caspase apoptosis are well known in the art and may be used or routinely modified to	Preferred embodiments of the invention include using polypeptides of the invention (or antibodies, agonists, or antagonists thereof) in detection, diagnosis, prevention,

and/or treatment of asthma, allergy, hypersensitivity and inflammation.	A highly preferred embodiment of the invention includes a method for stimulating natural killer cell
assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to regulate caspase protease-mediated apoptosis in immune cells (such as, for example, in mast cells). Mast cells are found in connective and mucosal tissues throughout the body, and their activation via immunoglobulin E-antigen, promoted by T helper cell type 2 cytokines, is an important component of allergic disease. Dysregulation of mast cell apoptosis may play a role in allergic disease and mast cell tumor survival. Exemplary assays for caspase apoptosis that may be used or routinely modified to test capase apoptosis activity induced by polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include the assays disclosed in: Masuda A, et al., J Biol Chem, 276(28):26107-26113 (2001); Yeatman CF 2nd, et al., J Exp Med, 192(8):1093-1103 (2000); Lee et al., Twomb 3(2): 75-80 (1996); the contents of each of which are herein incorporated by reference in its entirety. Immune cells that may be used according to these assays include mast cells such as the HMC becomercial sources). Exemplary immune cells that may be used according to these assays include mast cells such as the HMC become in the armond according to these assays include mast cells such as the HMC become in the may be used according to these assays include mast cells such as the HMC become in the may be used according to these assays include mast cells such as the HMC become in the may be used according to these	Kinase assay. Kinase assays, for example an Elk-1 kinase assay, for ERK signal
	Activation of Natural Killer Cell ERK
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	HADMB15
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	Signaling Pathway.	transduction that regulate cell proliferation	proliferation. An alternative highly preferred embodiment
		or differentiation are well known in the art	of the invention includes a method for inhibiting natural
		and may be used or routinely modified to	killer cell proliferation. A highly preferred
		assess the ability of polypeptides of the	embodiment of the invention includes a method for
		invention (including antibodies and	stimulating natural killer cell differentiation. An
		agonists or antagonists of the invention) to	alternative highly preferred embodiment of the invention
		promote or inhibit cell proliferation,	includes a method for inhibiting natural killer cell
		activation, and differentiation. Exemplary	differentiation. Highly preferred indications include
		assays for ERK kinase activity that may be	neoplastic diseases (e.g., as described below under
		used or routinely modified to test ERK	"Hyperproliferative Disorders"), blood disorders (e.g., as
		kinase-induced activity of polypeptides of	described below under "Immune Activity",
		the invention (including antibodies and	"Cardiovascular Disorders", and/or "Blood-Related
		agonists or antagonists of the invention)	Disorders"), immune disorders (e.g., as described below
		include the assays disclosed in Forrer et	under "Immune Activity") and infections (e.g., as
		al., Biol Chem 379(8-9):1101-1110	described below under "Infectious Disease"). Preferred
-		(1998); Kyriakis JM, Biochem Soc Symp	indications include blood disorders (e.g., as described
		64:29-48 (1999); Chang and Karin, Nature	below under "Immune Activity", "Blood-Related
		410(6824):37-40 (2001); and Cobb MH,	Disorders", and/or "Cardiovascular Disorders"). Highly
		Prog Biophys Mol Biol 71(3-4):479-500	preferred indications include autoimmune diseases (e.g.,
		(1999); the contents of each of which are	rheumatoid arthritis, systemic lupus erythematosis,
		herein incorporated by reference in its	multiple sclerosis and/or as described below) and
		entirety. Natural killer cells that may be	immunodeficiencies (e.g., as described below). Additional
		used according to these assays are publicly	highly preferred indications include inflammation and
		available (e.g., through the ATCC).	inflammatory disorders. Highly preferred indications
		Exemplary natural killer cells that may be	as,
		used according to these assays include the	breast, lung, colon, pancreatic, esophageal, stomach,
		human natural killer cell lines (for	brain, liver, urinary cancer, lymphoma and leukemias.
		example, NK-YT cells which have	Other preferred indications include benign dysproliferative
		cytolytic and cytotoxic activity) or primary	disorders and pre-neoplastic conditions, such as, for
		NK cells.	example, hyperplasia, metaplasia, and/or dysplasia.
			Other highly preferred indications include, pancytopenia,
			leukopenia, leukemias, Hodgkin's disease, acute
			lymphocytic anemia (ALL), arthritis, asthma, AIDS,
			granulomatous disease, inflammatory bowel disease,
			sepsis, psoriasis, immune reactions to transplanted organs
			and tissues, endocarditis, meningitis, Lyme Disease, and
			allergies.

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A highly preferred embodiment of the invention includes a method for stimulating the production of IFNg. An alternative highly preferred embodiment of the invention includes a method for inhibiting the production of IFNg. Highly preferred indications include blood disorders (e.g., as described below under "Immune Activity", "Blood-Related Disorders", and/or "Cardiovascular Disorders"), and infection (e.g., viral infections, tuberculosis, infections associated with chronic granulomatosus disease and malignant osteoporosis, and/or as described below under "Infectious Disease"). Highly preferred indications include autoimmune disease (e.g., rheumatoid arthritis, systemic lupus erythematosis, multiple sclerosis and/or as described below), boosting a Timmunodeficiency (e.g., as described below), boosting a T	cell-mediated immune response, and suppressing a T cell-mediated immune response. Additional highly preferred indications include inflammation and inflammatory disorders. Additional preferred indications include idiopathic pulmonary fibrosis. Highly preferred indications include neoplastic diseases (e.g., leukemia, lymphoma, melanoma, and/or as described below under "Hyperproliferative Disorders"). Highly preferred indications include neoplasms and cancers, such as, for example, leukemia, lymphoma, melanoma, and prostate, breast lung colon, parceasic secondarial stormach.	brain, lung, colon, pancleauc, esopuagea,, stornach, brain, liver and urinary cancer. Other preferred indications include benign dysproliferative disorders and preneoplastic conditions, such as, for example, hyperplasia, metaplasia, and/or dysplasia. Preferred indications include anemia, pancytopenia, leukopenia, thrombocytopenia, Hodgkin's disease, acute lymphocytic anemia (ALL), plasmacytomas, multiple myeloma, Burkitt's lymphoma, arthritis, AIDS, granulomatous disease, inflammatory bowel disease, sepsis, neutropenia, neutrophilia, psoriasis, suppression of immune reactions to
IFNgamma FMAT. IFNg plays a central role in the immune system and is considered to be a proinflammatory cytokine. IFNg promotes TH1 and inhibits TH2 differentiation; promotes IgG2a and inhibits IgE secretion; induces macrophage activation; and increases MHC expression. Assays for immunomodulatory proteins produced by T cells and NK cells that regulate a variety of inflammatory activities and inhibit TH2 helper cell functions are well known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and	agonists or antagonists of the invention) to mediate immunomodulation, regulate inflammatory activities, modulate TH2 helper cell function, and/or mediate humoral or cell-mediated immunity. Exemplary assays that test for immunomodulatory proteins evaluate the production of cytokines, such as Interferon gamma (IFNg), and the activation of T cells. Such assays that may be used or proteinely modified to test	immunomodulatory activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include the assays disclosed in Miraglia et al., J Biomolecular Screening 4:193-204 (1999); Rowland et al., "Lymphocytes: a practical approach" Chapter 6:138-160 (2000); Gonzalez et al., J Clin Lab Anal 8(5):225-233 (1995); Billiau et al., Ann NY Acad Sci 856:22-32 (1908). Brothm et al., Ann NY Acad Sci 856:22-32 (1908).
Production of IFNgamma using a T cells		
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hyr me	Highly preferred indications include blood disorders (e.g., as described below under "Immune Activity", "Blood-Related Disorders", and/or "Cardiovascular Disorders"). Highly preferred indications include autoimmune diseases (e.g., rheumatoid arthritis, systemic lupus erythematosis, multiple sclerosis and/or as described modulatory boosting a T cell-mediated immune response, and alternatively, suppressing a T cell-mediated immune response, and alternatively, suppressing a T cell-mediated immune response. A highly preferred indication is a complication associated with diabetes (e.g., diabetic neuropathy, higheric neuropathy, holod vessel blockage, heart disease, stroke, impotence (e.g., due to diabetic neuropathy, or blood vessel blockage), seizures, mental confusion, drowsiness, nonketotic hyperglycemic-says that may hyperosmolar coma, cardiovascular disease, and disorders as to test hypertension, stroke, and other diseases and disorders as
15:749-795 (1997), and Rheumatology (Oxford) 38(3):214-20 (1999), the contents of each of which are herein incorporated by reference in its entirety. Human T cells that may be used according to these assays may be isolated using techniques disclosed herein or otherwise known in the art. Human T cells are primary human lymphocytes that mature in the thymus and express a T Cell receptor and CD3, CD4, or CD8. These cells mediate humoral or cell-mediated immunity and may be preactivated to enhance responsiveness to immunomodulatory factors.	HLA-DR FMAT. MHC class II is essential for correct presentation of antigen to CD4+T cells. Deregulation of MHC class II has been associated with autoimmune diseases (e.g., diabetes, rheumatoid arthritis, systemic lupus erythematosis, and multiple sclerosis). Assays for immunomodulatory proteins expressed on MHC class II expressing T cells and antigen presenting cells are well known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to modulate the activation of T cells, and/or mediate humoral or cell-mediated immunity. Exemplary assays that test for immunomodulatory proteins evaluate the upregulation of MHC class II products, such as HLA-DR antigens, and the activation of T cells. Such assays that may be used or routinely modified to test immunomodulatory activity of
	Upregulation of HLA-DR and activation of T cells
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polypeptides of the invention (including	described in the "Cardiovascular Disorders" section
the invention) include, for example, the	the "Endocrine Disorders" section below), neuropathy,
assays disclosed in Miraglia et al., J	vision impairment (e.g., diabetic retinopathy and
Biomolecular Screening 4:193-204 (1999);	blindness), ulcers and impaired wound healing, and
Rowland et al., "Lymphocytes: a practical	infection (e.g., infectious diseases and disorders as
approach" Chapter 6:138-160 (2000);	described in the "Infectious Diseases" section below,
Lamour et al., Clin Exp Immunol	especially of the urinary tract and skin), carpal tunnel
89(2):217-222 (1992); Hurme and Sihvola,	syndrome and Dupuytren's contracture). An
Immunol Lett 20(3):217-222 (1989);	additional highly preferred indication is obesity and/or
Gansbacher and Zier, Cell Immunol	complications associated with obesity. Additional highly
117(1):22-34 (1988); and Itoh et al., J	preferred indications include weight loss or alternatively,
Histochem Cytochem 40(11):1675-1683,	weight gain. Aditional highly preferred indications
 the contents of each of which are herein	are complications associated with insulin resistance.
incorporated by reference in its entirety.	Additional highly preferred indications are disorders of the
Human T cells that may be used according	musculoskeletal systems including myopathies, muscular
to these assays may be isolated using	dystrophy, and/or as described herein.
techniques disclosed herein or otherwise	additional preferred indication is infection (e.g., AIDS,
known in the art. Human T cells are	and/or as described below under "Infectious Disease").
primary human lymphocytes that mature in	Preferred indications include endocrine disorders (e.g., as
the thymus and express a T Cell receptor	described below under "Endocrine Disorders"), and
and CD3, CD4, or CD8. These cells	neoplastic diseases (e.g., leukemia, lymphoma, and/or as
mediate humoral or cell-mediated	described below under "Hyperproliferative Disorders").
immunity and may be preactivated to	Preferred indications include neoplasms and cancer, such
 enhance responsiveness to	as, for example, leukemia, lymphoma, and prostate, breast,
 immunomodulatory factors.	lung, colon, pancreatic, esophageal, stomach, brain, liver
	and urinary cancer. Other preferred indications include
	benign dysproliferative disorders and pre-neoplastic
	conditions, such as, for example, hyperplasia, metaplasia,
	and/or dysplasia. Preferred indications also include
	anemia, pancytopenia, leukopenia, thrombocytopenia,
	Hodgkin's disease, acute lymphocytic anemia (ALL),
	plasmacytomas, multiple myeloma, Burkitt's lymphoma,
	arthritis, AIDS, granulomatous disease, inflammatory
	bowel disease, sepsis, neutropenia, neutrophilia, psoriasis,
 _	suppression of immune reactions to transplanted organs
	and tissues, nemopnilia, nypercoagulation, endocardius,

## COMPOST TOST

HAGEGIO 526 Activation of Assays for the activation of transcription through the Gamma Interferon Activation GAS response element in immune cells (such in immune cells (such coutinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention including antibodies and agonists or antagonists of the invention including antibodies and agonists or antagonists of the invention including cell functions. Exemplary assays for transcription factors and modulate gene expression involved in a wide variety of cell functions. Exemplary assays for transcription through the GAS response element activity of polypeptides of the invention (including antibodies and agonists or antagonists of antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthom et al., Proc Natl Acad Sci USA 85:632-6346 (1988); Matikainen et al., Blood 93(6):1990, the contents of each of which are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		asthma and allergy.
	Assays for the activation of transcription	Highly preferred indications include neoplastic diseases  o leukemia lymphoma and/or as described below
	Site (GAS) response element are well-	sorders"). Highly preferred
	known in the art and may be used or	is and cancers, such as, for
antibodies and agonists or antagonists of antibodies and agonists or antagonists of the invention) to regulate STAT transcription factors and modulate gene expression involved in a wide variety of cell functions. Exemplary assays for transcription through the GAS response element modified to test GAS-response element activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Matikalinen et al., Blood 93(6):1980-1991 (1999); and Henttinen et al., I Immunol 155(10):4582-4587 (1995), the contents of each of which are herein incoprorated by reference in its entirery. Exemplary mouse 7 cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTILL cell line, which is a suspension		na (e.g., T cell lymphoma,
antibodies and agonists or antagonists of the invention) to regulate STAT transcription factors and modulate gene expression involved in a wide variety of cell functions. Exemplary assays for transcription through the GAS response element that may be used or routinely modified to test GAS-response element activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988), Mattkainen et al., Blood 93(6):1880-1991 (1999); and Henttinen et al., J Immunol 155(10):4582-4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		Burkitt's lymphoma, non-Hodgkins lymphoma, Hodgkin's
transcription factors and modulate gene expression involved in a wide variety of cell functions. Exemplary assays for transcription through the GAS response element that may be used or routinely modified to test GAS-response element activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthom et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Matikainen et al., Blood 93(6):1980-1991 (1999); and Henttinen et al., Immunol 155(10):4882-4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell ine, which is a suspension	onists of	state, breast, lung, colon,
transcription factors and modulate gene expression involved in a wide variety of cell functions. Exemplary assays for transcription through the GAS response element that may be used or routinely modified to test GAS-response element activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998), Cullen and Malm, Methods in Enzymol 216:362-368 (1992), Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Matikainen et al., Blood 93(6):1980-1991 (1999); and Henttinen et al., J Immunol 155(10):4882-4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the CTL. Cell line, which is a suspension		ach, brain, liver and urinary
expression involved in a wide variety of cell functions. Exemplary assays for transcription through the GAS response element that may be used or routinely modified to test GAS-response element activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Matikainen et al., Blood 93(6):1980-1991 (1999); and Hentinen et al., Immunol 155(10):4582-4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		ations include benign
cell functions. Exemplary assays for transcription through the GAS response element that may be used or routinely modified to test GAS-response element activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1888); Matikainen et al., Blood 93(6):1980-1991 (1999); and Henttinen et al., I Immunol 155(10):4582-4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		d pre-neoplastic conditions,
transcription through the GAS response element that may be used or routinely modified to test GAS-response element activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Matikainen et al., Blood 93(6):1980-1991 (1999); and Henttinen et al., I Immunol 155(10):4582-4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		lasia, metaplasia, and/or
element that may be used or routinely modified to test GAS-response element activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Matikainen et al., Blood 93(6):1980-1991 (1999); and Hentinen et al., J Immunol 155(10):4582-4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		ations include autoimmune
modified to test GAS-response element activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Matikainen et al., Blood 93(6):1980-1991 (1999); and Henttinen et al., J Immunol 155(10):4582-4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		thritis, systemic lupus
activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthom et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Matikainen et al., Blood 93(6):1980-1991 (1999); and Henttinen et al., J Immunol 155(10):4582-4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		osis and/or as described
(including antibodies and agonists or antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Matikainen et al., Blood 93(6):1980-1991 (1999); and Henttinen et al., J Immunol 155(10):4582-4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		(e.g., as described below),
antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Matikainen et al., Blood 93(6):1980-1991 (1999); and Hentinen et al., J Immunol 155(10):4582-4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		mmune response, and
assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Matikainen et al., Blood 93(6):1980-1991 (1999); and Henttinen et al., J Immunol 155(10):4582-4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		d immune response.
66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Matikainen et al., Blood 93(6):1980-1991 (1999); and Hentinen et al., J Immunol 155(10):4582- 4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension	ene	Additional preferred indications include inflammation and
Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Matikainen et al., Blood 93(6):1980-1991 (1999); and Hentinen et al., J Immunol 155(10):4582-4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		Highly preferred indications
Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Matikainen et al., Blood 93(6):1980-1991 (1999); and Henttinne et al., J Immunol 155(10):4582-4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		, as described below under
85:6342-6346 (1988); Matikainen et al., Blood 93(6):1980-1991 (1999); and Henttinen et al., J Immunol 155(10):4582- 4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		Related Disorders", and/or
Blood 93(6):1980-1991 (1999); and Henttinen et al., J Immunol 155(10):4582- 4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		, and infection (e.g., viral
Henttinen et al., J Immunol 155(10):4582-4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		infections, tuberculosis, infections associated with chronic
4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		malignant osteoporosis,
are herein incorporated by reference in its entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		as described below under
entirety. Exemplary mouse T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension	ıts	onal
may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		osis. Preferred indications
publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		a, leukopenia,
ATCC). Exemplary T cells that may be used according to these assays include the CTLL cell line, which is a suspension		nphocytic anemia (ALL),
used according to these assays include the		eloma, arthritis, AIDS,
CTLL cell line, which is a suspension		mmatory bowel disease,
E :		sepsis, neutropenia, neutrophilia, psoriasis, suppression of
culture of LL-2 dependent cytotoxic i	culture of IL-2 dependent cytotoxic T immune reactions to transplanted organs and tissues,	inted organs and tissues,
cells.	cells. hemophilia, hypercoagulation, diabetes mellitus,	n, diabetes mellitus,

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13	HAGEQ79	527	Activation of	Assays for the activation of transcription	Highly preferred indications include neoplastic diseases
	,		transcription through	through the Gamma Interferon Activation	(e.g., leukemia, lymphoma, and/or as described below
			GAS response element	Site (GAS) response element are well-	under "Hyperproliferative Disorders"). Highly preferred
			in immune cells (such	known in the art and may be used or	indications include neoplasms and cancers, such as, for
			as T-cells).	routinely modified to assess the ability of	example, leukemia, lymphoma (e.g., T cell lymphoma,
				polypeptides of the invention (including	Burkitt's lymphoma, non-Hodgkins lymphoma, Hodgkin's
				antibodies and agonists or antagonists of	disease), melanoma, and prostate, breast, lung, colon,
				the invention) to regulate STAT	pancreatic, esophageal, stomach, brain, liver and urinary
				transcription factors and modulate gene	cancer. Other preferred indications include benign
				expression involved in a wide variety of	dysproliferative disorders and pre-neoplastic conditions,
				cell functions. Exemplary assays for	such as, for example, hyperplasia, metaplasia, and/or
				transcription through the GAS response	dysplasia. Preferred indications include autoimmune
				element that may be used or routinely	diseases (e.g., rheumatoid arthritis, systemic lupus
				modified to test GAS-response element	erythematosis, multiple sclerosis and/or as described
				activity of polypeptides of the invention	below), immunodeficiencies (e.g., as described below),
				(including antibodies and agonists or	boosting a T cell-mediated immune response, and
				antagonists of the invention) include	suppressing a T cell-mediated immune response.
				assays disclosed in Berger et al., Gene	Additional preferred indications include inflammation and
				66:1-10 (1998); Cullen and Malm,	inflammatory disorders. Highly preferred indications
				Methods in Enzymol 216:362-368 (1992);	include blood disorders (e.g., as described below under
				Henthorn et al., Proc Natl Acad Sci USA	"Immune Activity", "Blood-Related Disorders", and/or
				85:6342-6346 (1988); Matikainen et al.,	"Cardiovascular Disorders"), and infection (e.g., viral
				Blood 93(6):1980-1991 (1999); and	infections, tuberculosis, infections associated with chronic
				Henttinen et al., J Immunol 155(10):4582-	granulomatosus disease and malignant osteoporosis,
				4587 (1995), the contents of each of which	and/or an infectious disease as described below under
				are herein incorporated by reference in its	ınal
				entirety. Exemplary mouse T cells that	is idiopathic pulmonary fibrosis. Preferred indications
				may be used according to these assays are	include anemia, pancytopenia, leukopenia,
				publicly available (e.g., through the	thrombocytopenia, acute lymphocytic anemia (ALL),
				ATCC). Exemplary T cells that may be	plasmacytomas, multiple myeloma, arthritis, AIDS,
				used according to these assays include the	granulomatous disease, inflammatory bowel disease,
				CTLL cell line, which is a suspension	sepsis, neutropenia, neutrophilia, psoriasis, suppression of
				culture of IL-2 dependent cytotoxic T	immune reactions to transplanted organs and tissues,
				cells.	hemophilia, hypercoagulation, diabetes mellitus,
,					endocarditis, meningitis, Lyme Disease, and asthma and
					allergy.

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Highly preferred indications include blood disorders (e.g., as described below under "Immune Activity", "Blood-Related Disorders", and/or "Cardiovascular Disorders"). Highly preferred indications include autoimmune diseases (e.g., rheumatoid arthritis, systemic lupus erythematosis, multiple sclerosis and/or as described below).	boosting a T cell-mediated immune response, and alternatively, suppressing a T cell-mediated immune response.  A highly preferred indication is diabetes mellitus.  An additional highly preferred indication is a complication associated with diabetes (e.g., diabetic retinopathy, diabetic nephropathy, kidney diseases (e.g., renal failure, nephropathy and/or other diseases and disorders as described in the "Renal Disorders" section	below), diabetic neuropathy, nerve disease and nerve damage (e.g., due to diabetic neuropathy), blood vessel blockage, heart disease, stroke, impotence (e.g., due to diabetic neuropathy or blood vessel blockage), seizures, mental confusion, drowsiness, nonketotic hyperglycemichyperosmolar coma, cardiovascular disease (e.g., heart disease, atherosclerosis, microvascular disease, hypertension, stroke, and other diseases and disorders as described in the "Cardiovascular Disorders" section below), dystinidemia endocrine disorders (as described in	the "Endocrine Disorders" section below), neuropathy, vision impairment (e.g., diabetic retinopathy and blindness), ulcers and impaired wound healing, and infection (e.g., infectious diseases and disorders as described in the "Infectious Diseases" section below, especially of the urinary tract and skin), carpal tunnel syndrome and Dupuytren's contracture). An additional highly preferred indication is obesity and/or complications associated with obesity. Additional highly preferred indications include weight loss or alternatively, weight gain. Additional highly preferred indications associated with insulin resistance.
HLA-DR FMAT. MHC class II is essential for correct presentation of antigen to CD4+ T cells. Deregulation of MHC class II has been associated with autoimmune diseases (e.g., diabetes, rheumatoid arthritis, systemic lupus erythematosis, and multiple	scierosis). Assays for infinundulation y proteins expressed on MHC class II expressing T cells and antigen presenting cells are well known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to modulate the activation of T cells, and/or mediate	humoral or cell-mediated immunity.  Exemplary assays that test for immunomodulatory proteins evaluate the upregulation of MHC class II products, such as HLA-DR antigens, and the activation of T cells. Such assays that may be used or routinely modified to test immunomodulatory activity of polypeptides of the invention (including polypeptides of the invention (including polypeptides).	the invention) include, for example, the assays disclosed in Miraglia et al., J Biomolecular Screening 4:193-204 (1999); Rowland et al., "Lymphocytes: a practical approach" Chapter 6:138-160 (2000); Lamour et al., Clin Exp Immunol 89(2):217-222 (1992); Hurme and Sihvola, Immunol Lett 20(3):217-222 (1989); Gansbacher and Zier, Cell Immunol 117(1):22-34 (1988); and Itoh et al., J Histochem Cytochem 40(11):1675-1683, the contents of each of which are herein
Upregulation of HLA-DR and activation of T cells			
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ety. Additional highly preferred indications are disorders of the musculoskeletal systems including myopathies, muscular dystrophy, and/or as described herein. An additional preferred indication is infection (e.g., AIDS, and/or as described below under "Infectious Disease"). Preferred indications include endocrine disorders (e.g., as described below under "Hyperproliferative Disorders"). Preferred indications include neoplasms and cancer, such as, for example, leukemia, lymphoma, and/or as described below under "Hyperproliferative Disorders"). Preferred indications include neoplasms and cancer, such as, for example, leukemia, lymphoma, and prostate, breast, lung, colon, pancreatic, esophageal, stomach, brain, liver and urinary cancer. Other preferred indications include benign dysproliferative disorders and pre-neoplastic conditions, such as, for example, hyperplasia, metaplasia, and/or dysplasia. Preferred indications also include anemia, pancytopenia, leukopenia, thrombocytopenia, Hodgkin's disease, acute lymphocytic anemia (ALL), plasmacytomas, multiple myeloma, Burkitt's lymphoma, arthritis, AIDS, granulomatous disease, inflammatory bowel disease, sepsis, neutropenia, neutrophilia, psoriasis, suppression of immune reactions to transplanted organs and tissues, hemophilia, hypercoagulation, endocarditis, meningitis, Lyme Disease, inflammation and allergy.	s 4 , E
incorporated by reference in its entirety. Human T cells that may be used according to these assays may be isolated using techniques disclosed herein or otherwise known in the art. Human T cells are primary human lymphocytes that mature in the thymus and express a T Cell receptor and CD3, CD4, or CD8. These cells mediate humoral or cell-mediated immunity and may be preactivated to enhance responsiveness to immunomodulatory factors.	Kinase assays, for example kinase assays for members of the MAP kinase family (including p38, JAK, and ERK) are well known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to promote or inhibit cell proliferation, differentiation, and/or cytokine production in immune cells such as T-cells. Exemplary assays for MAP kinase family members that may be used
	Proliferation, differentiation, and/or cytokine production in immune cells (such as T-cells).
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				or routinely modified to test polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include the assays disclosed in: Rincon M., Curr Opin Immunol; 13(3):339-345 (2001); Wang LH, et al., J Immunol, 162(7):3897-3904 (1999); Sakamoto H, et al., J Biol Chem, 275(46):35857-35862 (2000), the contents of each of which are herein incorporated by reference in its	
				entirety. Exemplary immune cells (for example, T-cells) that may be used according to these assays include the mouse CTLL cell line.	
14	HAGFS57	528	Production of IL-6	IL-6 FMAT. IL-6 is produced by T cells and has strong effects on B cells. IL-6	A highly preferred embodiment of the invention includes a method for stimulating (e.g., increasing) IL-6
				participates in IL-4 induced IgE production and increases IgA production (IgA plays a	production. An alternative highly preferred embodiment of the invention includes a method for inhibiting (e.g.,
				role in mucosal immunity). IL-6 induces	reducing) IL-6 production. A highly preferrred
				cytotoxic T cells. Deregulated expression of II -6 has been linked to autoimmune	indication is the sumulation of enhancement of inucosal immunity. Highly preferred indications include blood
				disease, plasmacytomas, myelomas, and	disorders (e.g., as described below under "Immune
				chronic hyperproliferative diseases.	Activity", "Blood-Related Disorders", and/or
				Assays for immunomodulatory and	"Cardiovascular Disorders"), and infection (e.g., as
				differentiation factor proteins produced by	described below under injectious Disease ). riiginy preferred indications include autoimmune diseases (e.g.,
				expression level is strongly regulated by	rheumatoid arthritis, systemic lupus erythematosis,
				cytokines, growth factors, and hormones	multiple sclerosis and/or as described below) and imminodeficiencies (e.g., as described below). Highly
				or routinely modified to assess the ability	preferred indications also include boosting a B cell-
				of polypeptides of the invention (including	mediated immune response and alternatively suppressing a
				antibodies and agonists or antagonists of	B cell-mediated immune response. Highly preferred
				the invention) to mediate	indications include inflammation and inflammatory
				immunomodulation and differentiation and	disorders. Additional highly preferred indications include
				modulate T cell proliferation and function.	asthma and allergy. Highly preferred indications include
				Exemplary assays that test for	neoplastic diseases (e.g., myeloma, plasmacytoma,
				immunomodulatory proteins evaluate the	leukemia, lymphoma, melanoma, and/or as described

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				production of cytokines, such as IL-6, and	below under "Hyperproliferative Disorders"). Highly
				the stimulation and upregulation of T cell	preferred indications include neoplasms and cancers, such
				proliferation and functional activities.	as, myeloma, plasmacytoma, leukemia, lymphoma,
				Such assays that may be used or routinely	melanoma, and prostate, breast, lung, colon, pancreatic,
				modified to test immunomodulatory and	esophageal, stomach, brain, liver and urinary cancer.
				diffferentiation activity of polypeptides of	Other preferred indications include benign dysproliferative
				the invention (including antibodies and	disorders and pre-neoplastic conditions, such as, for
				agonists or antagonists of the invention)	example, hyperplasia, metaplasia, and/or dysplasia.
				include assays disclosed in Miraglia et al.,	Preferred indications include anemia, pancytopenia,
				J Biomolecular Screening 4:193-	leukopenia, thrombocytopenia, Hodgkin's disease, acute
				204(1999); Rowland et al., "Lymphocytes:	lymphocytic anemia (ALL), multiple myeloma, Burkitt's
				a practical approach" Chapter 6:138-160	lymphoma, arthritis, AIDS, granulomatous disease,
				(2000); and Verhasselt et al., J Immunol	inflammatory bowel disease, sepsis, neutropenia,
				158:2919-2925 (1997), the contents of	neutrophilia, psoriasis, suppression of immune reactions to
				each of which are herein incorporated by	transplanted organs and tissues, hemophilia,
				reference in its entirety. Human dendritic	hypercoagulation, diabetes mellitus, endocarditis,
				cells that may be used according to these	meningitis, and Lyme Disease. An additional preferred
				assays may be isolated using techniques	indication is infection (e.g., an infectious disease as
				disclosed herein or otherwise known in the	described below under "Infectious Disease").
				art. Human dendritic cells are antigen	
				presenting cells in suspension culture,	
				which, when activated by antigen and/or	
	-			cytokines, initiate and upregulate T cell	
				proliferation and functional activities.	
15	HAGHIN57	529	Activation of	Assays for the activation of transcription	A preferred embodiment of the invention includes a
			transcription through	through the Serum Response Element	method for inhibiting (e.g., reducing) TNF alpha
			serum response element	(SRE) are well-known in the art and may	production. An alternative highly preferred embodiment of
			in immune cells (such	be used or routinely modified to assess the	the invention includes a method for stimulating (e.g.,
			as T-cells).	ability of polypeptides of the invention	increasing) TNF alpha production. Preferred indications
				(including antibodies and agonists or	include blood disorders (e.g., as described below under
				antagonists of the invention) to bind the	"Immune Activity", "Blood-Related Disorders", and/or
				serum response factor and modulate the	"Cardiovascular Disorders"), Highly preferred indications
				expression of genes involved in growth	include autoimmune diseases (e.g., rheumatoid arthritis,
				and upregulate the function of growth-	systemic lupus erythematosis, Crohn's disease, multiple
				related genes in many cell types.	sclerosis and/or as described below), immunodeficiencies
				Exemplary assays for transcription through	(e.g., as described below), boosting a T cell-mediated
				the SRE that may be used or routinely	immune response, and suppressing a T cell-mediated

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				modified to test SRE activity of the polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Mathods in Engravol 216:362	immune response. Additional highly preferred indications include inflammation and inflammatory disorders, and treating joint damage in patients with rheumatoid arthritis. An additional highly preferred indication is sepsis. Highly preferred indications include neoplastic diseases
				Acad Sci USA 85:6342-6346 (1988); Benson et al., J Immunol 153(9):3862-3873 (1994); and Black et al., Virus Genes	under "Hyperproliferative Disorders"). Additionally, highly preferred indications include neoplasms and cancers, such as, for example, leukemia, lymphoma, melanoma, glioma (e.g., malignant glioma), solid tumors,
				of which are herein incorporated by reference in its entirety. T cells that may be used according to these assays are muhiicly available (e.g., through the	stomach, brain, liver and urinary cancer. Other preferred indications include benign dysproliferative disorders and pre-neoplastic conditions, such as, for example, hyperplasia metanlasia and/or dysplasia.
				ATCC). Exemplary human T cells, such as the MOLT4, that may be used according to these according	
				through the ATCC).	Burkitt's lymphoma, arthritis, AIDS, granulomatous disease, inflammatory bowel disease, neutropenia,
					neutrophilia, psoriasis, suppression or immune reactions to transplanted organs and tissues, hemophilia, hypercoagulation, diabetes mellitus, endocarditis,
					meningitis, Lyme Disease, cardiac reperfusion injury, and asthma and allergy. An additional preferred indication is infection (e.g., an infectious disease as described below index "Treorious Disease").
15	HAGHIN57	529	Activation of transcription through	Assays for the activation of transcription through the Signal Transducers and	A highly preferred indication is allergy.  Additional
			STAT6 response element in immune	Activators of Transcription (STAT6) response element are well-known in the art	nflammat erred indi
			cells (such as natural killer cells).	and may be used or routinely modified to assess the ability of polypeptides of the	"Imclude blood disorders (e.g., as described below under "Immune Activity", "Blood-Related Disorders", and/or
				invention (including antibodies and	"Cardiovascular Disorders"). Preferred indications include
				regulate STAT6 transcription factors and modulate the expression of multiple genes.	autoinment diseases (e.g., incumatory at firms, systeme lupus erythematosis, multiple sclerosis and/or as described below) and immunodeficiencies (e.g., as described below).

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				Exemplary assays for transcription through	Preferred indications include neoplastic diseases (e.g.,
				the STAT6 response element that may be	leukemia, lymphoma, melanoma, and/or as described
				used or routinely modified to test STAT6	below under "Hyperproliferative Disorders"). Preferred
				response element activity of the	indications include neoplasms, such as, for example,
				polypeptides of the invention (including	leukemia, lymphoma, melanoma, and prostate, breast,
				antibodies and agonists or antagonists of	lung, colon, pancreatic, esophageal, stomach, brain, liver
,				the invention) include assays disclosed in	and urinary cancer. Other preferred indications include
				Berger et al., Gene 66:1-10 (1998); Cullen	benign dysproliferative disorders and pre-neoplastic
				and Malm, Methods in Enzymol 216:362-	conditions, such as, for example, hyperplasia, metaplasia,
				368 (1992); Henthorn et al., Proc Natl	and/or dysplasia. Preferred indications include
				Acad Sci USA 85:6342-6346 (1988);	anemia, pancytopenia, leukopenia, thrombocytopenia,
				Georas et al., Blood 92(12):4529-4538	Hodgkin's disease, acute lymphocytic anemia (ALL),
				(1998); Moffatt et al., Transplantation	plasmacytomas, multiple myeloma, Burkitt's lymphoma,
				69(7):1521-1523 (2000); Curiel et al., Eur	arthritis, AIDS, granulomatous disease, inflammatory
-				J Immunol 27(8):1982-1987 (1997); and	bowel disease, sepsis, neutropenia, neutrophilia, psoriasis,
				Masuda et al., J Biol Chem	suppression of immune reactions to transplanted organs
				275(38):29331-29337 (2000), the contents	and tissues, hemophilia, hypercoagulation, diabetes
				of each of which are herein incorporated	mellitus, endocarditis, meningitis, and Lyme Disease.
				by reference in its entirety. T cells that	Additional preferred indications include infection (e.g., an
				may be used according to these assays are	infectious disease as described below under "Infectious
				publicly available (e.g., through the	Disease").
				ATCC). Exemplary rat natural killer cells	
_				that may be used according to these assays	
				are publicly available (e.g., through the	
				ATCC).	
15	HAGHIN57	529	Activation of	Assays for the activation of transcription	Highly preferred indications include blood disorders
			transcription through	through the Nuclear Factor of Activated T	(e.g., as described below under "Immune Activity",
			NFAT response	cells (NFAT) response element are well-	"Blood-Related Disorders", and/or "Cardiovascular
			element in immune	known in the art and may be used or	Disorders"). Highly preferred indications include
			cells (such as natural	routinely modified to assess the ability of	autoimmune diseases (e.g., rheumatoid arthritis, systemic
			killer cells).	polypeptides of the invention (including	lupus erythematosis, multiple sclerosis and/or as described
				antibodies and agonists or antagonists of	below), immunodeficiencies (e.g., as described below),
		,		the invention) to regulate NFAT	boosting a T cell-mediated immune response, and
				transcription factors and modulate	suppressing a T cell-mediated immune response.
				expression of genes involved in	Additional highly preferred indications include
				immunomodulatory functions. Exemplary	inflammation and inflammatory disorders. An additional
			7	assays for transcription through the NFAT	highly preferred indication is intection (e.g., an intectious

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disease as described below under "Infectious Disease"). Preferred indications include neoplastic diseases (e.g., leukemia, lymphoma, and/or as described below under "Hyperproliferative Disorders"). Preferred indications include neoplasms and cancers, such as, for example, leukemia, lymphoma, and prostate, breast, lung, colon, pancreatic, esophageal, stomach, brain, liver and urinary cancer. Other preferred indications include benign dysproliferative disorders and pre-neoplastic conditions, such as, for example, hyperplasia, metaplasia, and/or dysplasia. Preferred indications also include anemia, pancytopenia, leukopenia, thrombocytopenia, Hodgkin's disease, acute lymphocytic anemia (ALL), plasmacytomas, multiple myeloma, Burkitt's lymphoma, arthritis, AIDS, granulomatous disease, inflammatory bowel disease, sepsis, neutropenia, neutrophilia, psoriasis, suppression of immune reactions to transplanted organs and tissues, hemophilia, hypercoagulation, diabetes mellitus, endocarditis, meningitis, Lyme Disease, asthma and allergy.	A highly preferred embodiment of the invention includes a method for activating T cells. An alternative highly preferred embodiment of the invention includes a method for inhibiting the activation of and/or inactivating T cells. A highly preferred embodiment of the invention includes a method for activation B cells. An alternative highly preferred embodiment of the invention includes a method for inhibiting the activation of and/or inactivating B cells. A highly preferred embodiment of the invention includes a method for activating NK cells. An alternative highly preferred embodiment of the invention includes a method for inhibiting activation of
response element that may be used or routinely modified to test NFAT-response element activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Aramburu et al., J Exp Med 182(3):801-810 (1995); De Boer et al., Int J Biochem Cell Biol 31(10):1221-1236 (1999); Fraser et al., Eur J Immunol 29(3):838-844 (1999); and Yeseen et al., J Biol Chem 268(19):14285-14293 (1993), the contents of each of which are herein incorporated by reference in its entirety. NK cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary human NK cells that may be used according to these assays include the NK-YT cell line, which is a human natural killer cell line with cytolytic and cytotoxic activity.	CD69 FMAT. CD69 is an activation marker that is expressed on activated T cells, B cells, and NK cells. CD69 is not expressed on resting T cells, B cells, or NK cells. CD69 has been found to be associated with inflammation. Assays for immunomodulatory proteins expressed in T cells, B cells, and leukocytes are well known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of
	Upregulation of CD69 and activation of T cells
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			the invention) to modulate the activation of	and/or inactivation NK cells. Highly preferred
			T cells, and/or mediate humoral or cell-	indications include inflammation and inflammatory
			mediated immunity. Exemplary assays	disorders (e.g., as described below under "Immune Activity")
			evaluate the upregulation of cell surface	e.e.
			markers, such as CD69, and the activation	Activity", "Blood-Related Disorders", and/or
			of T cells. Such assays that may be used	"Cardiovascular Disorders"). Highly preferred indications
			or routinely modified to test	include autoimmune diseases (e.g., rheumatoid arthritis,
			immunomodulatory activity of	systemic lupus erythematosis, multiple sclerosis and/or as
			polypeptides of the invention (including	described below), immunodeficiencies (e.g., as described
			antibodies and agonists or antagonists of	below), boosting a T cell-mediated immune response and
			the invention) include, for example, the	alternatively suppressing a T cell-mediated immune
			assays disclosed in Miraglia et al., J	response, and boosting a B cell-mediated immune
			Biomolecular Screening 4:193-204 (1999);	response and alternatively suppressing a B cell-mediated
			Rowland et al., "Lymphocytes: a practical	immune response. An additional highly preferred
			approach" Chapter 6:138-160 (2000);	indication includes infection (e.g., as described below
			Ferenczi et al., J Autoimmun 14(1):63-78	under "Infectious Disease"). Preferred indications also
			(200); Werfel et al., Allergy 52(4):465-469	include anemia, pancytopenia, leukopenia,
			(1997); Taylor-Fishwick and Siegel, Eur J	thrombocytopenia, Hodgkin's disease, acute lymphocytic
			Immunol 25(12):3215-3221 (1995); and	anemia (ALL), plasmacytomas, multiple myeloma,
			Afetra et al., Ann Rheum Dis 52(6):457-	Burkitt's lymphoma, arthritis, AIDS, granulomatous
			460 (1993), the contents of each of which	disease, inflammatory bowel disease, sepsis, neutropenia,
			are herein incorporated by reference in its	neutrophilia, psoriasis, suppression of immune reactions to
			entirety. Human T cells that may be used	transplanted organs and tissues, hemophilia,
			according to these assays may be isolated	hypercoagulation, diabetes mellitus, endocarditis,
			using techniques disclosed herein or	meningitis, Lyme Disease, inflammation and
			otherwise known in the art. Human T cells	inflammatory disorders, asthma, and allergies.
			are primary human lymphocytes that	Preferred indications also include neoplastic diseases (e.g.,
			mature in the thymus and express a T Cell	leukemia, lymphoma, and/or as described below under
•			receptor and CD3, CD4, or CD8. These	"Hyperproliferative Disorders"). Preferred indications
			cells mediate humoral or cell-mediated	include neoplasms, such as, for example, leukemia,
			immunity and may be preactivated to	lymphoma, and prostate, breast, lung, colon, pancreatic,
			enhance responsiveness to	esophageal, stomach, brain, liver and urinary cancer.
			immunomodulatory factors.	Other preferred indications include benign dysproliferative
				disorders and pre-neoplastic conditions, such as, for
				example, hyperplasia, metaplasia, and/or dysplasia.
17 HAJAA47	531	Production of TNF	TNFa FMAT. Assays for	A highly preferred embodiment of the invention

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alpha by dend	intic cells activated macrophages, T smooth muscle, and other exert a wide variety of inf cytotoxic effects on a variwell known in the art and routinely modified to asse polypeptides of the inventantibodies and agonists of the inventantibodies and agonists of the inventantibodies and agonists of the invention) to mediate immunomodulation, modification and cytotox assays that test for immun proteins evaluate the prodiction of an inflamma response. Such assays the routinely modified to test immunomodulatory activity polypeptides of the inventantibodies and agonists of the inventantibodies agonists of the inventantibodies agonists of the inventantibodies agonists and agonists and agonists of the inventantibodies agonises and agonises and agonises and agonises and agonises and agon	immunomodulatory proteins produced by activated macrophages, T cells, fibroblasts, smooth muscle, and other cell types that exert a wide variety of inflammatory and cytotoxic effects on a variety of cells are well known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to mediate imflammation and cytotoxicity. Exemplary assays that test for immunomodulatory proteins evaluate the production of cytokines such as tumor necrosis factor alpha (TNFa), and the induction or inhibition of an inflammatory or cytotoxic response. Such assays that may be used or routinely modified to test immunomodulatory activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in Miraglia et al., J Biomolecular Screening 4:193-204(1999); Rowland et al., "Lymphocytes: a practical approach"	includes a method for inhibiting (e.g., decreasing) TNF alpha production. An alternative highly preferred embodiment of the invention includes a method for stimulating (e.g., increasing) TNF alpha production. Highly preferred indications include blood disorders (e.g., as described below under "Immune Activity", "Blood-Related Disorders", and/or "Cardiovascular Disorders"), Highly preferred indications include autoimmune diseases (e.g., rheumatoid arthritis, systemic lupus erythematosis, Crohn's disease, multiple sclerosis and/or as described below), boosting a T cell-mediated immune response. Additional highly preferred indications include inflammation and inflammatory disorders, and treating joint damage in patients with rheumatoid arthritis. An additional highly preferred indication is sepsis. Highly preferred indications include neoplastic diseases (e.g., leukemia, lymphoma, and/or as described below under "Hyperproliferative Disorders"). Additionally, highly preferred indications include neoplasms and cancers, such as, leukemia, lymphoma, melanoma, glioma (e.g., malignant glioma), solid tumors, and prostate, breast, lung, colon, pancreatic, esophageal, stomach, brain, liver and urinary cancer. Other preferred indications include benign dysproliferative disorders and pre-neoplastic conditions such as for example by hyperplasia
	al., Eur J Imm (1198); Dahle (107):3585-3 al., J Immunol Nardelli et al., (1999), the co herein incorpc entirety. Hum be used accorr isolated using	Chapter 0:138-100 (2000); Vernassell et al., Eur J Immunol 28(11):3886-3890 (1198); Dahlen et al., J Immunol 160(7):3585-3593 (1998); Verhasselt et al., J Immunol 158:2919-2925 (1997); and Nardelli et al., J Leukoc Biol 65:822-828 (1999), the contents of each of which are herein incorporated by reference in its entirety. Human dendritic cells that may be used according to these assays may be isolated using techniques disclosed herein	conditions, such as, for example, hyperplassa, metaplassa, and/or dysplassa. Preferred indications include anemia, pancytopenia, leukopenia, thrombocytopenia, Hodgkin's disease, acute lymphocytic anemia (ALL), plasmacytomas, multiple myeloma, Burkitt's lymphoma, arthritis, AIDS, granulomatous disease, inflammatory bowel disease, neutropenia, neutrophilia, psoriasis, suppression of immune reactions to transplanted organs and tissues, hemophilia, hypercoagulation, diabetes mellitus, endocarditis, meningitis, Lyme Disease, cardiac reperfusion injury, and asthma and allergy. An

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HAJAA47	531	Activation of transcription through the EGR (Early Growth Response) element in immune cells (such as B-cells).	dendritic cells are antigen presenting cells in suspension culture, which, when activated by antigen and/or cytokines, initiate and upregulate T cell proliferation and functional activities.  Assays for the activation of transcription through the EGR response element are well-known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to regulate EGR transcription factors and modulate expression of immunomodulatory genes. Exemplary assays for transcription through the EGR response element that may be used or routinely modified to test EGR response element activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in: Richards JD, include assays disclosed in: Richards JD,	infectious disease as described below under "Infectious Disease").  Preferred embodiments of the invention include using polypeptides of the invention (or antibodies, agonists, or antagonists thereof) in detection, diagnosis, prevention, and/or treatment of Cancer, Autoimmunity, Allergy and Asthma.
HAJAŸ92	532	Activation of transcription through GAS response element	et al., J Immunol, 166(6):3855-3864 (2001); Dinkel, A, et al., J Exp Med, 188(12):2215-2224 (1998); and, Newton, JS, et al., Eur J Immunol 1996 Apr;26(4):811-816 (1996), the contents of each of which are herein incorporated by reference in its entirety. Immune cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary immune cells that may be used according to these assays include the Raji B-cell line.  Assays for the activation of transcription through the Gamma Interferon Activation Site (GAS) response element are well-	Preferred embodiments of the invention include using polypeptides of the invention (or antibodies, agonists, or antagonists thereof) in detection, diagnosis, prevention,

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and/or treatment of Inflammation, Infection, Cancer, Hypersensitivity, and Atherosclerosis.  J  S:  S:  S:  S:  S:  S:  S:  S:  S:		A highly preferred indication is diabetes mellitus.  An additional highly preferred indication is a complication associated with diabetes (e.g., diabetic retinopathy,
known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to regulate STAT transcription factors and modulate gene expression involved in a wide variety of cell functions. Exemplary assays for transcription through the GAS response element activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in: Gustafson KS, et al., J Biol Chem, 271(33):20035-20046 (1996); Eilers A, et al., Immunobiology, 193(2-4):328-333 (1995); Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Matikainen et al., Blood 93(6):1980-1991 (1999); and Henttinen et al., J Immunol 155(10):4582-4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary immune cells that may be used according to these assays are publicly available (e.g., through the	may be used according to these assays include the U937 cell line, which is a monocytic cell line.	Assays for measuring secretion of insulin are well-known in the art and may be used or routinely modified to assess the ability of notwestides of the invantion (including
in immune cells (such as monocytes).		Stimulation of insulin secretion from pancreatic beta cells.
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